

# Katherine Belov

## 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.

193  
papers

5,741  
citations

36  
h-index

66  
g-index

207  
ext. papers

6,724  
ext. citations

5.4  
avg, IF

5.58  
L-index

#	Paper	IF	Citations
193	Threatened Species Initiative: Empowering conservation action using genomic resources.. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2022</b> , 119,	11.5	8
192	Future-proofing the koala: synergising genomic and environmental data for effective species management.. <i>Molecular Ecology</i> , <b>2022</b> ,	5.7	4
191	Incomplete lineage sorting and phenotypic evolution in marsupials.. <i>Cell</i> , <b>2022</b> ,	56.2	5
190	Epigenetic clock and methylation studies in marsupials: opossums, Tasmanian devils, kangaroos, and wallabies.. <i>GeroScience</i> , <b>2022</b> , 1	8.9	0
189	A targeted approach to investigating immune genes of an iconic Australian marsupial.. <i>Molecular Ecology</i> , <b>2022</b> ,	5.7	3
188	How much is enough? Sampling intensity influences estimates of reproductive variance in an introduced population. <i>Ecological Applications</i> , <b>2021</b> , e02462	4.9	1
187	Metapopulation management of a critically endangered marsupial in the age of genomics. <i>Global Ecology and Conservation</i> , <b>2021</b> , 31, e01869	2.8	3
186	Koala cathelicidin PhciCath5 has antimicrobial activity, including against <i>Chlamydia pecorum</i> . <i>PLoS ONE</i> , <b>2021</b> , 16, e0249658	3.7	1
185	First evidence of deviation from Mendelian proportions in a conservation programme. <i>Molecular Ecology</i> , <b>2021</b> , 30, 3703-3715	5.7	1
184	Characterization of reproductive gene diversity in the endangered Tasmanian devil. <i>Molecular Ecology Resources</i> , <b>2021</b> , 21, 721-732	8.4	0
183	Assessing evolutionary processes over time in a conservation breeding program: a combined approach using molecular data, simulations and pedigree analysis. <i>Biodiversity and Conservation</i> , <b>2021</b> , 30, 1011-1029	3.4	2
182	Annotation of immune genes in the extinct thylacine ( <i>Thylacinus cynocephalus</i> ). <i>Immunogenetics</i> , <b>2021</b> , 73, 263-275	3.2	1
181	Improved high-throughput MHC typing for non-model species using long-read sequencing. <i>Molecular Ecology Resources</i> , <b>2021</b> ,	8.4	1
180	Platypus and echidna genomes reveal mammalian biology and evolution. <i>Nature</i> , <b>2021</b> , 592, 756-762	50.4	28
179	Comprehensive Knowledge of Reservoir Hosts is Key to Mitigating Future Pandemics. <i>Innovation(China)</i> , <b>2020</b> , 1, 100065	17.8	1
178	Marsupial Gut Microbiome. <i>Frontiers in Microbiology</i> , <b>2020</b> , 11, 1058	5.7	0
177	A demonstration of conservation genomics for threatened species management. <i>Molecular Ecology Resources</i> , <b>2020</b> , 20, 1526-1541	8.4	23

176	Inbreeding depression in one of the last DFTD-free wild populations of Tasmanian devils. <i>PeerJ</i> , <b>2020</b> , 8, e9220	3.1	1
175	Genetic analysis of scat samples to inform conservation of the Tasmanian devil. <i>Australian Zoologist</i> , <b>2020</b> , 40, 492-504	0.7	
174	Deciphering genetic mate choice: Not so simple in group-housed conservation breeding programs. <i>Evolutionary Applications</i> , <b>2020</b> , 13, 2179-2189	4.8	3
173	Too much of a good thing? Finding the most informative genetic data set to answer conservation questions. <i>Molecular Ecology Resources</i> , <b>2019</b> , 19, 659-671	8.4	18
172	From reference genomes to population genomics: comparing three reference-aligned reduced-representation sequencing pipelines in two wildlife species. <i>BMC Genomics</i> , <b>2019</b> , 20, 453	4.5	24
171	Tasmanian devils with contagious cancer exhibit a constricted T-cell repertoire diversity. <i>Communications Biology</i> , <b>2019</b> , 2, 99	6.7	11
170	Fecal Viral Diversity of Captive and Wild Tasmanian Devils Characterized Using Virion-Enriched Metagenomics and Metatranscriptomics. <i>Journal of Virology</i> , <b>2019</b> , 93,	6.6	29
169	MHC-associated mate choice under competitive conditions in captive versus wild Tasmanian devils. <i>Behavioral Ecology</i> , <b>2019</b> , 30, 1196-1204	2.3	4
168	Molecular characterisation of Interleukin-2 in two Australian marsupials (the tammar wallaby, <i>Notamacropus eugenii</i> , and the Tasmanian devil, <i>Sarcophilus harrisii</i> ) facilitates the development of marsupial-specific immunological reagents. <i>Australian Mammalogy</i> , <b>2019</b> , 41, 39	1.1	1
167	A Tasmanian devil breeding program to support wild recovery. <i>Reproduction, Fertility and Development</i> , <b>2019</b> , 31, 1296-1304	1.8	14
166	Looking like the locals - gut microbiome changes post-release in an endangered species. <i>Animal Microbiome</i> , <b>2019</b> , 1, 8	4.1	21
165	The Value of Reference Genomes in the Conservation of Threatened Species. <i>Genes</i> , <b>2019</b> , 10,	4.2	36
164	Tasmanian Devil Facial Tumor Disease <b>2019</b> , 490-493		
163	Complex problems need detailed solutions: Harnessing multiple data types to inform genetic management in the wild. <i>Evolutionary Applications</i> , <b>2019</b> , 12, 280-291	4.8	20
162	Koala immunology and infectious diseases: How much can the koala bear?. <i>Developmental and Comparative Immunology</i> , <b>2018</b> , 82, 177-185	3.2	5
161	Gomesin peptides prevent proliferation and lead to the cell death of devil facial tumour disease cells. <i>Cell Death Discovery</i> , <b>2018</b> , 4, 19	6.9	10
160	Transcriptomic changes in the pre-implantation uterus highlight histotrophic nutrition of the developing marsupial embryo. <i>Scientific Reports</i> , <b>2018</b> , 8, 2412	4.9	14
159	Landscape-level field data reveal broad-scale effects of a fatal, transmissible cancer on population ecology of the Tasmanian devil. <i>Mammalian Biology</i> , <b>2018</b> , 91, 41-45	1.6	6

158	MHC diversity and female age underpin reproductive success in an Australian icon; the Tasmanian Devil. <i>Scientific Reports</i> , <b>2018</b> , 8, 4175	4.9	13
157	Characterisation of MHC class I genes in the koala. <i>Immunogenetics</i> , <b>2018</b> , 70, 125-133	3.2	12
156	Pedigree reconstruction using molecular data reveals an early warning sign of gene diversity loss in an island population of Tasmanian devils ( <i>Sarcophilus harrisii</i> ). <i>Conservation Genetics</i> , <b>2018</b> , 19, 439-450	2.6	21
155	Devil women. <i>Pacific Conservation Biology</i> , <b>2018</b> , 24, 271	1.2	
154	Immunization Strategies Producing a Humoral IgG Immune Response against Devil Facial Tumor Disease in the Majority of Tasmanian Devils Destined for Wild Release. <i>Frontiers in Immunology</i> , <b>2018</b> , 9, 259	8.4	26
153	The effects of group versus intensive housing on the retention of genetic diversity in insurance populations. <i>BMC Zoology</i> , <b>2018</b> , 3,	1.8	17
152	Bioluminescent murine models of bacterial sepsis and scald wound infections for antimicrobial efficacy testing. <i>PLoS ONE</i> , <b>2018</b> , 13, e0200195	3.7	16
151	Disentangling the mechanisms of mate choice in a captive koala population. <i>PeerJ</i> , <b>2018</b> , 6, e5438	3.1	4
150	Marsupial Immunology <b>2018</b> ,		1
149	Women in conservation science making a difference. <i>Pacific Conservation Biology</i> , <b>2018</b> , 24, 209	1.2	7
148	Lessons learnt from the Tasmanian devil facial tumour regarding immune function in cancer. <i>Mammalian Genome</i> , <b>2018</b> , 29, 731-738	3.2	4
147	Adaptation and conservation insights from the koala genome. <i>Nature Genetics</i> , <b>2018</b> , 50, 1102-1111	36.3	102
146	Comparative genomics of hormonal signaling in the chorioallantoic membrane of oviparous and viviparous amniotes. <i>General and Comparative Endocrinology</i> , <b>2017</b> , 244, 19-29	3	24
145	Immune-endocrine interactions in marsupials and monotremes. <i>General and Comparative Endocrinology</i> , <b>2017</b> , 244, 178-185	3	5
144	Regression of devil facial tumour disease following immunotherapy in immunised Tasmanian devils. <i>Scientific Reports</i> , <b>2017</b> , 7, 43827	4.9	42
143	Variants in the host genome may inhibit tumour growth in devil facial tumours: evidence from genome-wide association. <i>Scientific Reports</i> , <b>2017</b> , 7, 423	4.9	38
142	The Regulation of Uterine Proinflammatory Gene Expression during Pregnancy in the Live-Bearing Lizard, <i>Pseudemoia entrecasteauxii</i> . <i>Journal of Experimental Zoology Part B: Molecular and Developmental Evolution</i> , <b>2017</b> , 328, 334-346	1.8	7
141	Significant decline in anticancer immune capacity during puberty in the Tasmanian devil. <i>Scientific Reports</i> , <b>2017</b> , 7, 44716	4.9	18

140	Long-read genome sequence assembly provides insight into ongoing retroviral invasion of the koala germline. <i>Scientific Reports</i> , <b>2017</b> , 7, 15838	4.9	27
139	No evidence of inbreeding depression in a Tasmanian devil insurance population despite significant variation in inbreeding. <i>Scientific Reports</i> , <b>2017</b> , 7, 1830	4.9	26
138	Increasing generations in captivity is associated with increased vulnerability of Tasmanian devils to vehicle strike following release to the wild. <i>Scientific Reports</i> , <b>2017</b> , 7, 2161	4.9	31
137	Characterization of the antimicrobial peptide family defensins in the Tasmanian devil ( <i>Sarcophilus harrisii</i> ), koala ( <i>Phascolarctos cinereus</i> ), and tammar wallaby ( <i>Macropus eugenii</i> ). <i>Immunogenetics</i> , <b>2017</b> , 69, 133-143	3.2	8
136	Devil Tools & Tech Synergy of Conservation Research and Management Practice. <i>Conservation Letters</i> , <b>2017</b> , 10, 133-138	6.9	26
135	Variation in Major Histocompatibility Complex diversity in invasive cane toad populations. <i>Wildlife Research</i> , <b>2017</b> , 44, 565	1.8	2
134	Antimicrobial Protection of Marsupial Pouch Young. <i>Frontiers in Microbiology</i> , <b>2017</b> , 8, 354	5.7	9
133	Marsupial and monotreme cathelicidins display antimicrobial activity, including against methicillin-resistant <i>Staphylococcus aureus</i> . <i>Microbiology (United Kingdom)</i> , <b>2017</b> , 163, 1457-1465	2.9	8
132	Reptile Pregnancy Is Underpinned by Complex Changes in Uterine Gene Expression: A Comparative Analysis of the Uterine Transcriptome in Viviparous and Oviparous Lizards. <i>Genome Biology and Evolution</i> , <b>2016</b> , 8, 3226-3239	3.9	29
131	Characterisation of the immune compounds in koala milk using a combined transcriptomic and proteomic approach. <i>Scientific Reports</i> , <b>2016</b> , 6, 35011	4.9	16
130	Allelic expression of mammalian imprinted genes in a matrotrophic lizard, <i>Pseudemoia entrecasteauxii</i> . <i>Development Genes and Evolution</i> , <b>2016</b> , 226, 79-85	1.8	14
129	The Platypus: A Venomous Mammal <b>2016</b> , 169-183		3
128	The identification of immune genes in the milk transcriptome of the Tasmanian devil ( <i>Sarcophilus harrisii</i> ). <i>PeerJ</i> , <b>2016</b> , 4, e1569	3.1	13
127	The Immune System of Monotremes and Marsupials <b>2016</b> , 504-514		
126	Evolution and comparative analysis of the bat MHC-I region. <i>Scientific Reports</i> , <b>2016</b> , 6, 21256	4.9	29
125	Immunoglobulin dynamics and cancer prevalence in Tasmanian devils ( <i>Sarcophilus harrisii</i> ). <i>Scientific Reports</i> , <b>2016</b> , 6, 25093	4.9	12
124	Molecular characterization of MHC class II in the Australian invasive cane toad reveals multiple splice variants. <i>Immunogenetics</i> , <b>2016</b> , 68, 449-460	3.2	7
123	Transmissible cancers in an evolutionary context. <i>BioEssays</i> , <b>2016</b> , 38 Suppl 1, S14-23	4.1	18

122	Transmissible cancers in an evolutionary context. <i>Inside the Cell</i> , <b>2016</b> , 1, 17-26		1
121	Diversity in the Toll-like receptor genes of the Tasmanian devil ( <i>Sarcophilus harrisii</i> ). <i>Immunogenetics</i> , <b>2015</b> , 67, 195-201	3.2	20
120	Cancer Immunology of Transmissible Cancers <b>2015</b> , 419-428		
119	Selection on MHC class II supertypes in the New Zealand endemic Hochstetter's frog. <i>BMC Evolutionary Biology</i> , <b>2015</b> , 15, 63	3	23
118	Impacts of early viability selection on management of inbreeding and genetic diversity in conservation. <i>Molecular Ecology</i> , <b>2015</b> , 24, 1645-53	5.7	13
117	Characterization of antibody V segment diversity in the Tasmanian devil ( <i>Sarcophilus harrisii</i> ). <i>Veterinary Immunology and Immunopathology</i> , <b>2015</b> , 167, 156-65	2	7
116	Identification and analysis of divergent immune gene families within the Tasmanian devil genome. <i>BMC Genomics</i> , <b>2015</b> , 16, 1017	4.5	10
115	Lack of genetic diversity across diverse immune genes in an endangered mammal, the Tasmanian devil ( <i>Sarcophilus harrisii</i> ). <i>Molecular Ecology</i> , <b>2015</b> , 24, 3860-72	5.7	43
114	Evolution of the avian $\beta$ -defensin and cathelicidin genes. <i>BMC Evolutionary Biology</i> , <b>2015</b> , 15, 188	3	47
113	Development of a SNP-based assay for measuring genetic diversity in the Tasmanian devil insurance population. <i>BMC Genomics</i> , <b>2015</b> , 16, 791	4.5	28
112	Identification, characterisation and expression analysis of natural killer receptor genes in <i>Chlamydia pecorum</i> infected koalas ( <i>Phascolarctos cinereus</i> ). <i>BMC Genomics</i> , <b>2015</b> , 16, 796	4.5	9
111	The Tasmanian devil microbiome-implications for conservation and management. <i>Microbiome</i> , <b>2015</b> , 3, 76	16.6	84
110	The origin, dynamics, and molecular evolution of transmissible cancers. <i>Advances in Genomics and Genetics</i> , <b>2015</b> , 317		
109	SNP marker discovery in koala TLR genes. <i>PLoS ONE</i> , <b>2015</b> , 10, e0121068	3.7	5
108	Genomic insights into a contagious cancer in Tasmanian devils. <i>Trends in Genetics</i> , <b>2015</b> , 31, 528-35	8.5	21
107	Anthropogenic selection enhances cancer evolution in Tasmanian devil tumours. <i>Evolutionary Applications</i> , <b>2014</b> , 7, 260-5	4.8	19
106	The koala immunological toolkit: sequence identification and comparison of key markers of the koala ( <i>Phascolarctos cinereus</i> ) immune response. <i>Australian Journal of Zoology</i> , <b>2014</b> , 62, 195	0.5	15
105	Characterisation of major histocompatibility complex class I in the Australian cane toad, <i>Rhinella marina</i> . <i>PLoS ONE</i> , <b>2014</b> , 9, e102824	3.7	18

104	Characterisation of non-classical MHC class I genes in the Tasmanian devil ( <i>Sarcophilus harrisii</i> ). <i>Immunogenetics</i> , <b>2014</b> , 66, 727-35	3.2	23
103	Identification of dendritic cells, B cell and T cell subsets in Tasmanian devil lymphoid tissue; evidence for poor immune cell infiltration into devil facial tumors. <i>Anatomical Record</i> , <b>2014</b> , 297, 925-38	2.1	29
102	Development of MHC-Linked Microsatellite Markers in the Domestic Cat and Their Use to Evaluate MHC Diversity in Domestic Cats, Cheetahs, and Gir Lions. <i>Journal of Heredity</i> , <b>2014</b> , 105, 493-505	2.4	7
101	Tracing monotreme venom evolution in the genomics era. <i>Toxins</i> , <b>2014</b> , 6, 1260-73	4.9	9
100	The Platypus: A Venomous Mammal <b>2014</b> , 1-13		
99	Reversible epigenetic down-regulation of MHC molecules by devil facial tumour disease illustrates immune escape by a contagious cancer. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2013</b> , 110, 5103-8	11.5	158
98	Low major histocompatibility complex diversity in the Tasmanian devil predates European settlement and may explain susceptibility to disease epidemics. <i>Biology Letters</i> , <b>2013</b> , 9, 20120900	3.6	40
97	Identification of natural killer cell receptor genes in the genome of the marsupial Tasmanian devil ( <i>Sarcophilus harrisii</i> ). <i>Immunogenetics</i> , <b>2013</b> , 65, 25-35	3.2	20
96	Characterisation of four major histocompatibility complex class II genes of the koala ( <i>Phascolarctos cinereus</i> ). <i>Immunogenetics</i> , <b>2013</b> , 65, 37-46	3.2	22
95	Does the devil facial tumour produce immunosuppressive cytokines as an immune evasion strategy?. <i>Veterinary Immunology and Immunopathology</i> , <b>2013</b> , 153, 159-64	2	19
94	Evolution of a contagious cancer: epigenetic variation in Devil Facial Tumour Disease. <i>Proceedings of the Royal Society B: Biological Sciences</i> , <b>2013</b> , 280, 20121720	4.4	17
93	Placental lipoprotein lipase (LPL) gene expression in a placentotrophic lizard, <i>Pseudemoia entrecasteauxii</i> . <i>Journal of Experimental Zoology Part B: Molecular and Developmental Evolution</i> , <b>2013</b> , 320, 465-70	1.8	8
92	Desmond Wishart Cooper: a life in science. <i>Australian Journal of Zoology</i> , <b>2013</b> , 61, 1	0.5	
91	Marsupial immunology bounding ahead. <i>Australian Journal of Zoology</i> , <b>2013</b> , 61, 24	0.5	17
90	Echidna venom gland transcriptome provides insights into the evolution of monotreme venom. <i>PLoS ONE</i> , <b>2013</b> , 8, e79092	3.7	14
89	Immune escape strategies of a contagious cancer, devil facial tumour disease. <i>Molecular Immunology</i> , <b>2012</b> , 51, 30	4.3	2
88	Contagious cancer: lessons from the devil and the dog. <i>BioEssays</i> , <b>2012</b> , 34, 285-92	4.1	27
87	Evolution in a transmissible cancer: a study of the chromosomal changes in devil facial tumor (DFT) as it spreads through the wild Tasmanian devil population. <i>Cancer Genetics</i> , <b>2012</b> , 205, 101-12	2.3	60



86	Molecular identification of interleukin-2 in the lymphoid tissues of the common brushtail possum, <i>Trichosurus vulpecula</i> . <i>Developmental and Comparative Immunology</i> , <b>2012</b> , 36, 236-40	3.2	9
85	Venom evolution through gene duplications. <i>Gene</i> , <b>2012</b> , 496, 1-7	3.8	73
84	Diversity at the major histocompatibility complex Class II in the platypus, <i>Ornithorhynchus anatinus</i> . <i>Journal of Heredity</i> , <b>2012</b> , 103, 467-78	2.4	7
83	Antigen-presenting genes and genomic copy number variations in the Tasmanian devil MHC. <i>BMC Genomics</i> , <b>2012</b> , 13, 87	4.5	47
82	Telomere dynamics and homeostasis in a transmissible cancer. <i>PLoS ONE</i> , <b>2012</b> , 7, e44085	3.7	19
81	A comparative genomics approach to understanding transmissible cancer in Tasmanian devils. <i>Annual Review of Genomics and Human Genetics</i> , <b>2012</b> , 13, 207-22	9.7	15
80	Isolation and characterisation of 11 MHC-linked microsatellite loci in the Tasmanian devil ( <i>Sarcophilus harrisii</i> ). <i>Conservation Genetics Resources</i> , <b>2012</b> , 4, 463-465	0.8	13
79	Low MHC class II diversity in the Tasmanian devil ( <i>Sarcophilus harrisii</i> ). <i>Immunogenetics</i> , <b>2012</b> , 64, 525-33	3.2	52
78	Reduced effect of Tasmanian devil facial tumor disease at the disease front. <i>Conservation Biology</i> , <b>2012</b> , 26, 124-34	6	58
77	A limited role for gene duplications in the evolution of platypus venom. <i>Molecular Biology and Evolution</i> , <b>2012</b> , 29, 167-77	8.3	30
76	Genomic restructuring in the Tasmanian devil facial tumour: chromosome painting and gene mapping provide clues to evolution of a transmissible tumour. <i>PLoS Genetics</i> , <b>2012</b> , 8, e1002483	6	75
75	Proteomics and deep sequencing comparison of seasonally active venom glands in the platypus reveals novel venom peptides and distinct expression profiles. <i>Molecular and Cellular Proteomics</i> , <b>2012</b> , 11, 1354-64	7.6	35
74	Diversity of MHC class II DAB1 in the koala ( <i>Phascolarctos cinereus</i> ). <i>Australian Journal of Zoology</i> , <b>2012</b> , 60, 1	0.5	13
73	New insights into the role of MHC diversity in devil facial tumour disease. <i>PLoS ONE</i> , <b>2012</b> , 7, e36955	3.7	28
72	Allorecognition in the Tasmanian devil ( <i>Sarcophilus harrisii</i> ), an endangered marsupial species with limited genetic diversity. <i>PLoS ONE</i> , <b>2011</b> , 6, e22402	3.7	55
71	Genome sequence of an Australian kangaroo, <i>Macropus eugenii</i> , provides insight into the evolution of mammalian reproduction and development <b>2011</b> , 12, 414		18
70	Genome sequence of an Australian kangaroo, <i>Macropus eugenii</i> , provides insight into the evolution of mammalian reproduction and development. <i>Genome Biology</i> , <b>2011</b> , 12, R81	18.3	142
69	Ancient antimicrobial peptides kill antibiotic-resistant pathogens: Australian mammals provide new options. <i>PLoS ONE</i> , <b>2011</b> , 6, e24030	3.7	61



68	The role of the Major Histocompatibility Complex in the spread of contagious cancers. <i>Mammalian Genome</i> , <b>2011</b> , 22, 83-90	3.2	40
67	Transcriptomic analysis supports similar functional roles for the two thymuses of the tammar wallaby. <i>BMC Genomics</i> , <b>2011</b> , 12, 420	4.5	19
66	The tammar wallaby major histocompatibility complex shows evidence of past genomic instability. <i>BMC Genomics</i> , <b>2011</b> , 12, 421	4.5	31
65	Immunome database for marsupials and monotremes. <i>BMC Immunology</i> , <b>2011</b> , 12, 48	3.7	22
64	Genomic identification of chemokines and cytokines in opossum. <i>Journal of Interferon and Cytokine Research</i> , <b>2011</b> , 31, 317-30	3.5	9
63	Major Histocompatibility Complex (MHC) markers in conservation biology. <i>International Journal of Molecular Sciences</i> , <b>2011</b> , 12, 5168-86	6.3	73
62	MHC gene copy number variation in Tasmanian devils: implications for the spread of a contagious cancer. <i>Proceedings of the Royal Society B: Biological Sciences</i> , <b>2010</b> , 277, 2001-6	4.4	104
61	Cloning and structural analysis of two highly divergent IgA isotypes, IgA1 and IgA2 from the duck billed platypus, <i>Ornithorhynchus anatinus</i> . <i>Molecular Immunology</i> , <b>2010</b> , 47, 785-91	4.3	11
60	Novel venom gene discovery in the platypus. <i>Genome Biology</i> , <b>2010</b> , 11, R95	18.3	45
59	L-to-D-peptide isomerase in male echidna venom. <i>Australian Journal of Zoology</i> , <b>2010</b> , 58, 284	0.5	11
58	Evolution of viviparity and uterine angiogenesis: vascular endothelial growth factor (VEGF) in oviparous and viviparous skinks. <i>Journal of Experimental Zoology Part B: Molecular and Developmental Evolution</i> , <b>2010</b> , 314, 148-56	1.8	9
57	MHC screening for marsupial conservation: extremely low levels of class II diversity indicate population vulnerability for an endangered Australian marsupial. <i>Conservation Genetics</i> , <b>2010</b> , 11, 269-278 <sup>6</sup>		13
56	The Marsupial Major Histocompatibility Complex <b>2010</b> , 339-356		1
55	Use of Genomic Information to Gain Insights into Immune Function in Marsupials: A Review of Divergent Immune Genes <b>2010</b> , 381-400		4
54	Hatching time for monotreme immunology. <i>Australian Journal of Zoology</i> , <b>2009</b> , 57, 185	0.5	9
53	Physical mapping of immune genes in the tammar wallaby ( <i>Macropus eugenii</i> ). <i>Cytogenetic and Genome Research</i> , <b>2009</b> , 127, 21-5	1.9	4
52	MHC-linked and un-linked class I genes in the wallaby. <i>BMC Genomics</i> , <b>2009</b> , 10, 310	4.5	40
51	High levels of genetic variation at MHC class II DBB loci in the tammar wallaby ( <i>Macropus eugenii</i> ). <i>Immunogenetics</i> , <b>2009</b> , 61, 111-8	3.2	17

50	Comparative genomics indicates the mammalian CD33rSiglec locus evolved by an ancient large-scale inverse duplication and suggests all Siglecs share a common ancestral region. <i>Immunogenetics</i> , <b>2009</b> , 61, 401-17	3.2	34
49	Identification of natural killer cell receptor clusters in the platypus genome reveals an expansion of C-type lectin genes. <i>Immunogenetics</i> , <b>2009</b> , 61, 565-79	3.2	22
48	Understanding and utilising mammalian venom via a platypus venom transcriptome. <i>Journal of Proteomics</i> , <b>2009</b> , 72, 155-64	3.9	32
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34	Transmission of a fatal clonal tumor by biting occurs due to depleted MHC diversity in a threatened carnivorous marsupial. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2007</b> , 104, 16221-6	11.5	219
33	Platypus Venom: a Review. <i>Australian Mammalogy</i> , <b>2007</b> , 29, 57	1.1	19

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