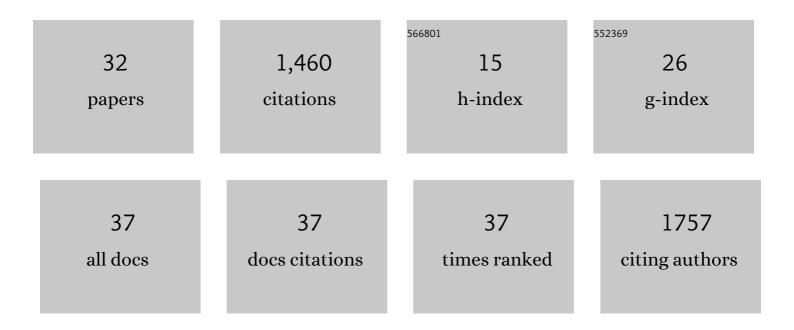
## Stuart J E Baird

List of Publications by Year in descending order

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STUADT I F RAIDO

#	Article	IF	CITATIONS
1	Hybridization, introgression, and linkage evolution. Plant Molecular Biology, 2000, 42, 205-224.	2.0	194
2	Rapid hybrid speciation in wild sunflowers. Proceedings of the National Academy of Sciences of the United States of America, 1998, 95, 11757-11762.	3.3	178
3	A Comparison of Multilocus Clines Maintained by Environmental Adaptation or by Selection Against Hybrids. Genetics, 1999, 153, 1959-1971.	1.2	170
4	ASSESSING MULTILOCUS INTROGRESSION PATTERNS: A CASE STUDY ON THE MOUSE X CHROMOSOME IN CENTRAL EUROPE. Evolution; International Journal of Organic Evolution, 2011, 65, 1428-1446.	1.1	108
5	Genetic conflict outweighs heterogametic incompatibility in the mouse hybrid zone?. BMC Evolutionary Biology, 2008, 8, 271.	3.2	94
6	PATTERNS OF MATING IN WILD SUNFLOWER HYBRID ZONES. Evolution; International Journal of Organic Evolution, 1998, 52, 713-726.	1.1	75
7	Sperm-related phenotypes implicated in both maintenance and breakdown of a natural species barrier in the house mouse. Proceedings of the Royal Society B: Biological Sciences, 2012, 279, 4803-4810.	1.2	60
8	Combining genetic, historical and geographical data to reconstruct the dynamics of bioinvasions: application to the cane toad <i>Bufo marinus</i> . Molecular Ecology Resources, 2010, 10, 886-901.	2.2	54
9	When Viruses Don't Go Viral: The Importance of Host Phylogeographic Structure in the Spatial Spread of Arenaviruses. PLoS Pathogens, 2017, 13, e1006073.	2.1	52
10	The complex social environment of female house mice ( <i>Mus domesticus</i> ). , 2012, , 114-134.		47
11	WHERE ARE THE WORMY MICE? A REEXAMINATION OF HYBRID PARASITISM IN THE EUROPEAN HOUSE MOUSE HYBRID ZONE. Evolution; International Journal of Organic Evolution, 2012, 66, 2757-2772.	1.1	47
12	The mouse hybrid zone in Central Europe: from morphology to molecules. Folia Zoologica, 2012, 61, 308-318.	0.9	41
13	What can the <i>Mus musculus musculus/M. m. domesticus</i> hybrid zone tell us about speciation?. , 2012, , 334-372.		37
14	On the origin of the house mouse synanthropy and dispersal in the Near East and Europe:. , 2012, , 65-93.		37
15	Empirical evidence for large X-effects in animals with undifferentiated sex chromosomes. Scientific Reports, 2016, 6, 21029.	1.6	35
16	Exploring linkage disequilibrium. Molecular Ecology Resources, 2015, 15, 1017-1019.	2.2	28
17	Genetic structure and contrasting selection pattern at two major histocompatibility complex genes in wild house mouse populations. Heredity, 2011, 106, 727-740.	1.2	27
18	Hybrid male sterility genes in the mouse subspecific crosses. , 2012, , 482-503.		23

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#	Article	IF	CITATIONS
19	Holobiont suture zones: Parasite evidence across the European house mouse hybrid zone. Molecular Ecology, 2018, 27, 5214-5227.	2.0	18
20	Murine Cytomegalovirus Is Not Restricted to the House Mouse Mus musculus domesticus: Prevalence and Genetic Diversity in the European House Mouse Hybrid Zone. Journal of Virology, 2015, 89, 406-414.	1.5	16
21	Genetic distinction between contiguous urban and rural multimammate mice in Tanzania despite gene flow. Journal of Evolutionary Biology, 2016, 29, 1952-1967.	0.8	14
22	Testing parasite â€`intimacy': the whipworm <i><scp>T</scp>richuris muris</i> inÂthe <scp>E</scp> uropean house mouse hybrid zone. Ecology and Evolution, 2016, 6, 2688-2701.	0.8	14
23	Host subspecific viral strains in European house mice: Murine cytomegalovirus in the Eastern (Mus) Tj ETQq1 1 (	0.784314 1.1	rgBT/Overloo
24	Intensity of infection with intracellular <i>Eimeria</i> spp. and pinworms is reduced in hybrid mice compared to parental subspecies. Journal of Evolutionary Biology, 2020, 33, 435-448.	0.8	11
25	The impact of global selection on local adaptation and reproductive isolation. Philosophical Transactions of the Royal Society B: Biological Sciences, 2020, 375, 20190531.	1.8	11
26	Bat population recoveries give insight into clustering strategies during hibernation. Frontiers in Zoology, 2020, 17, 26.	0.9	11
27	New insights into parasitism in the house mouse hybrid zone. , 2012, , 455-481.		9
28	The impact of highâ€ŧhroughput sequencing technology on speciation research: maintaining perspective. Journal of Evolutionary Biology, 2017, 30, 1482-1487.	0.8	9
29	Monte Carlo integration over stepping stone models for spatial genetic inference using approximate Bayesian computation. Molecular Ecology Resources, 2010, 10, 873-885.	2.2	8
30	Shifting Paradigms for Studying Parasitism in Hybridising Hosts: Response to Theodosopoulos, Hund, and Taylor. Trends in Ecology and Evolution, 2019, 34, 387-389.	4.2	7
31	A dense linkage map for a large repetitive genome: discovery of the sex-determining region in hybridizing fire-bellied toads ( <i>Bombina bombina</i> and <i>Bombina variegata</i> ). G3: Genes, Genomes, Genetics, 2021, 11, .	0.8	2
32	Evolutionary Ecology: Next Generation Inference. Current Biology, 2012, 22, R182-R183.	1.8	0