## H R Siegismund

List of Publications by Year in descending order

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471509 713466 1,378 22 17 21 citations h-index g-index papers 22 22 22 2529 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Mountain gorilla genomes reveal the impact of long-term population decline and inbreeding. Science, 2015, 348, 242-245.	12.6	326
2	Challenges for Serology-Based Characterization of Foot-and-Mouth Disease Outbreaks in Endemic Areas; Identification of Two Separate Lineages of Serotype O FMDV in Uganda in 2011. Transboundary and Emerging Diseases, 2015, 62, 522-534.	3.0	15
3	Understanding geographic origins and history of admixture among chimpanzees in European zoos, with implications for future breeding programmes. Heredity, 2013, 110, 586-593.	2.6	39
4	Extensive X-linked adaptive evolution in central chimpanzees. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 2054-2059.	7.1	79
5	Comparative phylogeography of African savannah ungulates < sup > 1 < /sup > . Molecular Ecology, 2012, 21, 3656-3670.	3.9	197
6	Molecular characterization of SAT 2 foot-and-mouth disease virus from post-outbreak slaughtered animals: implications for disease control in Uganda. Epidemiology and Infection, 2010, 138, 1204-1210.	2.1	19
7	Co-circulation of two extremely divergent serotype SAT 2 lineages in Kenya highlights challenges to foot-and-mouth disease control. Archives of Virology, 2010, 155, 1625-1630.	2.1	18
8	Relationship between three measures of genetic differentiation <i>G</i> <sub>ST</sub> , <i>D</i> <sub>EST</sub> and <i>G</i> ' <sub>ST</sub> : how wrong have we been?. Molecular Ecology, 2009, 18, 2080-2083.	3.9	151
9	Effective population size dynamics reveal impacts of historic climatic events and recent anthropogenic pressure in African elephants. Molecular Ecology, 2008, 17, 3788-3799.	3.9	47
10	Midâ€Holocene decline in African buffalos inferred from Bayesian coalescentâ€based analyses of microsatellites and mitochondrial DNA. Molecular Ecology, 2008, 17, 4845-4858.	3.9	50
11	Population Genetic Structure of Savannah Elephants in Kenya: Conservation and Management Implications. Journal of Heredity, 2008, 99, 443-452.	2.4	33
12	Mitochondrial DNA variation of the common hippopotamus: evidence for a recent population expansion. Heredity, 2005, 95, 206-215.	2.6	33
13	Noninvasive Genotyping and Mendelian Analysis of Microsatellites in African Savannah Elephants. Journal of Heredity, 2005, 96, 679-687.	2.4	25
14	Phylogeography and population structure of the common warthog (Phacochoerus africanus) inferred from variation in mitochondrial DNA sequences and microsatellite loci. Heredity, 2003, 91, 361-372.	2.6	74
15	Disparity in Population Differentiation of Sex-Linked and Autosomal Variation in Sibling Species of the Jaera albifrons (Isopoda) Complex., 2002, 93, 432-438.		4
16	Population structure of the African savannah elephant inferred from mitochondrial control region sequences and nuclear microsatellite loci. Heredity, 2002, 89, 90-98.	2.6	65
17	Population structure of African buffalo inferred from mtDNA sequences and microsatellite loci: high variation but low differentiation. Molecular Ecology, 1998, 7, 225-237.	3.9	89
18	Genetic structure and parasite compatibility of Bulinus globosus (Gastropoda: Planorbidae) from two areas of different endemicity of Schistosoma haematobium in Zimbabwe. International Journal for Parasitology, 1996, 26, 269-280.	3.1	18

#	Article	IF	CITATION
19	Mother-offspring data in a study of the mating system in a natural population of Bulinus globosus (Gastropoda: Planorbidae) in Zimbabwe. Genetical Research, 1996, 68, 95-100.	0.9	3
20	Extreme genetic differences among populations of Gazella granti, Grant's gazelle, in Kenya. Heredity, 1996, 76, 465-475.	2.6	73
21	Structure of African Elephant Populations. Journal of Heredity, 1995, 86, 467-469.	2.4	10
22	A Sex-Linked Enzyme Polymorphism in the Marine Isopod Jaera ischiosetosa. Journal of Heredity, 1992, 83, 388-393.	2.4	10