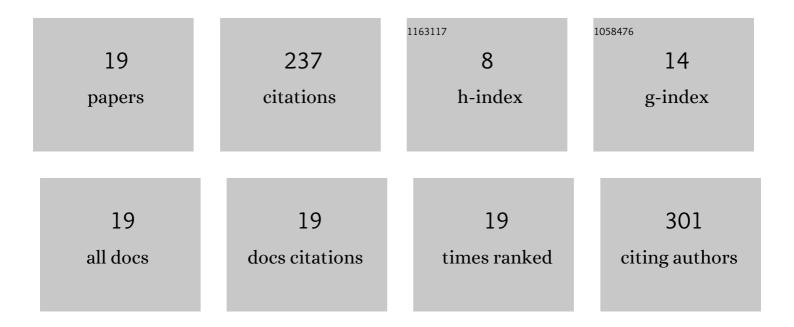
Xin-Sheng Hu

List of Publications by Year in descending order

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XIN-SHENC HU

#	Article	IF	CITATIONS
1	Impacts of Seed and Pollen Flow on Population Genetic Structure for Plant Genomes With Three Contrasting Modes of Inheritance. Genetics, 1999, 152, 441-450.	2.9	43
2	Mating system as a barrier to gene flow. Evolution; International Journal of Organic Evolution, 2015, 69, 1158-1177.	2.3	33
3	Screening Reliable Reference Genes for RT-qPCR Analysis of Gene Expression in Moringa oleifera. PLoS ONE, 2016, 11, e0159458.	2.5	22
4	Background selection and population differentiation. Journal of Theoretical Biology, 2005, 235, 207-219.	1.7	20
5	Seed and pollen flow in expanding a species' range. Journal of Theoretical Biology, 2006, 240, 662-672.	1.7	19
6	On migration load of seeds and pollen grains in a local population. Heredity, 2003, 90, 162-168.	2.6	15
7	Phylogeographic pattern suggests a general northeastward dispersal in the distribution of Machilus pauhoi in South China. PLoS ONE, 2017, 12, e0184456.	2.5	11
8	Mating system and population structure in the natural distribution of Toona ciliata (Meliaceae) in South China. Scientific Reports, 2020, 10, 16998.	3.3	11
9	BARRIERS TO THE SPREAD OF NEUTRAL ALLELES IN THE CYTONUCLEAR SYSTEM. Evolution; International Journal of Organic Evolution, 2008, 62, 2260-2278.	2.3	9
10	Estimating the number of sexual events per generation in a facultatively sexual haploid population. Heredity, 2019, 122, 729-741.	2.6	8
11	Structural Genomics: Correlation Blocks, Population Structure, and Genome Architecture. Current Genomics, 2011, 12, 55-70.	1.6	7
12	Mating system and the critical migration rate for swamping selection. Genetical Research, 2011, 93, 233-254.	0.9	6
13	High mutation rates explain low population genetic divergence at copy-number-variable loci in Homo sapiens. Scientific Reports, 2017, 7, 43178.	3.3	6
14	Advances in the study of population genetic diversity at plant species' margins. Chinese Journal of Plant Ecology, 2019, 43, 383-395.	0.6	6
15	A complete sequence of mitochondrial genome of Neolamarckia cadamba and its use for systematic analysis. Scientific Reports, 2021, 11, 21452.	3.3	6
16	Genomic Scans of Zygotic Disequilibrium and Epistatic SNPs in HapMap Phase III Populations. PLoS ONE, 2015, 10, e0131039.	2.5	5
17	Testing neutrality at copy-number-variable loci under the finite-allele and finite-site models. Theoretical Population Biology, 2016, 112, 1-13.	1.1	4
18	Mating system shifts a species' range. Evolution; International Journal of Organic Evolution, 2019, 73, 158-174.	2.3	4

#	Article	IF	CITATIONS
19	The wave of gene advance under diverse systems of mating. Heredity, 2020, 125, 253-268.	2.6	2