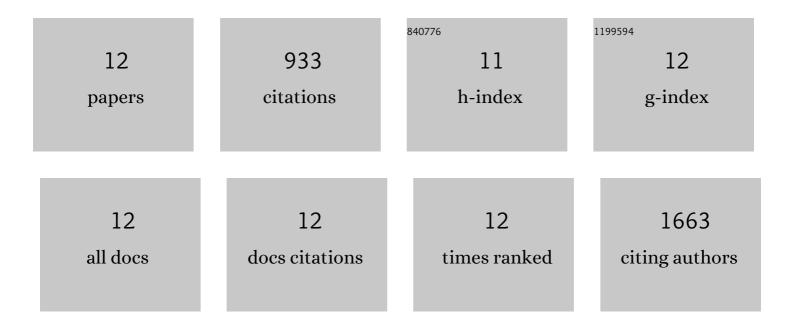
## **Mahesh Panchal**

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7982781/publications.pdf Version: 2024-02-01



Μλήξεη Ρληςήλι

#	Article	IF	CITATIONS
1	THE AUTOMATION AND EVALUATION OF NESTED CLADE PHYLOGEOGRAPHIC ANALYSIS. Evolution; International Journal of Organic Evolution, 2007, 61, 1466-1480.	2.3	143
2	In defence of model-based inference in phylogeography. Molecular Ecology, 2010, 19, 436-446.	3.9	141
3	Genomics of Divergence along a Continuum of Parapatric Population Differentiation. PLoS Genetics, 2015, 11, e1004966.	3.5	135
4	The automation of Nested Clade Phylogeographic Analysis. Bioinformatics, 2007, 23, 509-510.	4.1	127
5	Ten steps to get started in Genome Assembly and Annotation. F1000Research, 2018, 7, 148.	1.6	85
6	Genomeâ€wide patterns of standing genetic variation in a marine population of threeâ€spined sticklebacks. Molecular Ecology, 2013, 22, 635-649.	3.9	78
7	Extensive Copy-Number Variation of Young Genes across Stickleback Populations. PLoS Genetics, 2014, 10, e1004830.	3.5	70
8	On the validity of nested clade phylogeographical analysis. Molecular Ecology, 2008, 17, 2563-2565.	3.9	49
9	Transcriptome profiling of immune tissues reveals habitatâ€specific gene expression between lake and river sticklebacks. Molecular Ecology, 2016, 25, 943-958.	3.9	49
10	Evaluating Nested Clade Phylogeographic Analysis under Models of Restricted Gene Flow. Systematic Biology, 2010, 59, 415-432.	5.6	30
11	Identification of candidate mimicry proteins involved in parasite-driven phenotypic changes. Parasites and Vectors, 2015, 8, 225.	2.5	20
12	Differential Expression of Immune Genes between Two Closely Related Beetle Species with Different Immunocompetence following Attack by Asecodes parviclava. Genome Biology and Evolution, 2020, 12, 522-534.	2.5	6