Sergey V Malov

List of Publications by Citations

25 papers	**537** citations	**8** h-index	**23** g-index
26 ext. papers	**678** ext. citations	**3.7** avg, IF	**3.62** L-index

#	Paper	IF	Citations
25	The Genome 10K Project: a way forward. *Annual Review of Animal Biosciences*, **2015**, 3, 57-111	13.7	223
24	141 Gene Discovery and Data Sharing in Genome Wide Association Analyses. *Journal of Acquired Immune Deficiency Syndromes (1999)*, **2014**, 65, 59	3.1	78
23	The principal genetic determinants for nasopharyngeal carcinoma in China involve the HLA class I antigen recognition groove. *PLoS Genetics*, **2012**, 8, e1003103	6	72
22	Sympatric Asian felid phylogeography reveals a major Indochinese-Sundaic divergence. *Molecular Ecology*, **2014**, 23, 2072-92	5.7	38
21	Emerging viruses in the Felidae: shifting paradigms. *Viruses*, **2012**, 4, 236-57	6.2	37
20	A peer-educator network HIV prevention intervention among injection drug users: results of a randomized controlled trial in St. Petersburg, Russia. *AIDS and Behavior*, **2013**, 17, 2510-20	4.3	29
19	The novelty of human cancer/testis antigen encoding genes in evolution. *International Journal of Genomics*, **2013**, 2013, 105108	2.5	16
18	HIV incidence and behavioral correlates of HIV acquisition in a cohort of injection drug users in St Petersburg, Russia. *Medicine (United States)*, **2016**, 95, e5238	1.8	13
17	Genome-wide sequence analyses of ethnic populations across Russia. *Genomics*, **2020**, 112, 442-458	4.3	8
16	Genome-Wide Analyses Reveal Gene Influence on HIV Disease Progression and HIV-1C Acquisition in Southern Africa. *AIDS Research and Human Retroviruses*, **2017**, 33, 597-609	1.6	7
15	Semiparametric Regression Estimation in Copula Models. *Communications in Statistics - Theory and Methods*, **2006**, 35, 1449-1467	0.5	4
14	A Beautiful Life: High Risk-High Payoff in Genetic Science. *Annual Review of Animal Biosciences*, **2020**, 8, 1-24	13.7	3
13	GWATCH: a web platform for automated gene association discovery analysis. *GigaScience*, **2014**, 3, 18	7.6	2
12	Nonparametric estimation for a current status right-censored data model. *Statistica Neerlandica*, **2019**, 73, 475-495	0.9	1
11	Homogeneity Tests for Related Samples Under Censoring. *Communications in Statistics Part B: Simulation and Computation*, **2012**, 41, 764-775	0.6	1
10	Random variables generated by ranks in dependent schemes. *Metrika*, **1998**, 48, 61-67	0.8	1
9	Semiparametric Estimation in Copulas with the Same Marginals. *Communications in Statistics - Theory and Methods*, **2008**, 38, 372-392	0.5	1

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8	Testing of the homogeneity of marginal distributions in copula models. *Comptes Rendus Mathematique*, **2008**, 346, 445-450	0.4	1
7	On characterizations of independent identically distributed random variables via order structures. *Statistics and Probability Letters*, **1996**, 26, 97-104	0.6	1
6	Life table estimator revisited. *Communications in Statistics - Theory and Methods*, **2018**, 47, 2126-2133	0.5	1
5	Signal localization: a new approach in signal discovery. *Biometrical Journal*, **2017**, 59, 126-144	1.5	0
4	Evolution: a new cat species emerges. *Current Biology*, **2013**, 23, R1103-5	6.3	
3	Copula Process, Archimedean **2015**, 1-7		
2	Asymptotic Properties of Generalized Multivariate Rank Statistics. *Communications in Statistics - Theory and Methods*, **2012**, 41, 2133-2159	0.5	
1	Uniform convergence rate of the nonparametric maximum likelihood estimator for current status data with competing risks. *Statistics*, **2021**, 55, 152-172	0.5	