

# Hervé Perdry

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1230473/publications.pdf>

Version: 2024-02-01

17  
papers

274  
citations

1163117

8  
h-index

940533

16  
g-index

19  
all docs

19  
docs citations

19  
times ranked

395  
citing authors

#	ARTICLE	IF	CITATIONS
1	Estimating cancer risk in HNPCC by the GRL method. <i>European Journal of Human Genetics</i> , 2007, 15, 831-836.	2.8	54
2	The hCOMET project: International database comparison of results with the comet assay in human biomonitoring. Baseline frequency of DNA damage and effect of main confounders. <i>Mutation Research - Reviews in Mutation Research</i> , 2021, 787, 108371.	5.5	45
3	DNA damage in circulating leukocytes measured with the comet assay may predict the risk of death. <i>Scientific Reports</i> , 2021, 11, 16793.	3.3	36
4	Collection and storage of human white blood cells for analysis of DNA damage and repair activity using the comet assay in molecular epidemiology studies. <i>Mutagenesis</i> , 2021, 36, 193-212.	2.6	20
5	Accuracy of heritability estimations in presence of hidden population stratification. <i>Scientific Reports</i> , 2016, 6, 26471.	3.3	19
6	The Use of the Linear Mixed Model in Human Genetics. <i>Human Heredity</i> , 2015, 80, 196-206.	0.8	18
7	First genome-wide association study of non-severe malaria in two birth cohorts in Benin. <i>Human Genetics</i> , 2019, 138, 1341-1357.	3.8	14
8	Validation of Gelbond <sup>®</sup> high-throughput alkaline and Fpg <sup>®</sup> modified comet assay using a linear mixed model. <i>Environmental and Molecular Mutagenesis</i> , 2018, 59, 595-602.	2.2	13
9	The ordered transmission disequilibrium test: detection of modifier genes. <i>Genetic Epidemiology</i> , 2009, 33, 1-5.	1.3	9
10	Rare variant association testing for multicategory phenotype. <i>Genetic Epidemiology</i> , 2019, 43, 646-656.	1.3	9
11	An ordered subset approach to including covariates in the transmission disequilibrium test. <i>BMC Proceedings</i> , 2007, 1, S77.	1.6	8
12	Estimating penetrance from multiple case families with predisposing mutations: extension of the $\tilde{\alpha}$ -genotype-restricted likelihood <sup>™</sup> (GRL) method. <i>European Journal of Human Genetics</i> , 2011, 19, 173-179.	2.8	8
13	Adipose Tissue Properties in Tumor-Bearing Breasts. <i>Frontiers in Oncology</i> , 2020, 10, 1506.	2.8	6
14	A new scoring system in cancer genetics: application to criteria for BRCA1 and BRCA2 mutation screening. <i>Journal of Medical Genetics</i> , 2014, 51, 114-121.	3.2	4
15	Extension of SKAT to multi-category phenotypes through a geometrical interpretation. <i>European Journal of Human Genetics</i> , 2021, 29, 736-744.	2.8	4
16	Detecting the dominance component of heritability in isolated and outbred human populations. <i>Scientific Reports</i> , 2018, 8, 18048.	3.3	3
17	Mixed logistic regression in genome-wide association studies. <i>BMC Bioinformatics</i> , 2020, 21, 536.	2.6	3