

Penelope R Haddrill

List of Publications by Year in descending order

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33
papers

1,955
citations

361045

20
h-index

395343

33
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33
all docs

33
docs citations

33
times ranked

2369
citing authors

#	ARTICLE	IF	CITATIONS
1	Research trends in forensic science: A scientometric approach to analyze the content of the <scp>INTERPOL</scp> reviews. Wiley Interdisciplinary Reviews Forensic Science, 2022, 4, e1447.	1.2	5
2	Developments in forensic DNA analysis. Emerging Topics in Life Sciences, 2021, 5, 381-393.	1.1	36
3	Population genetics and forensic utility of 23 autosomal PowerPlex Fusion 6C STR loci in the Kuwaiti population. Scientific Reports, 2021, 11, 1865.	1.6	6
4	Population genetics of 30 insertion/deletion polymorphisms in the Kuwaiti population. International Journal of Legal Medicine, 2020, 134, 985-986.	1.2	5
5	Evaluating the effect of body fluid mixture on the relative expression ratio of blood-specific RNA markers. Forensic Science International, 2020, 307, 110116.	1.3	9
6	Determination of the most effective enhancement process for latent fingermarks on Clydesdale Bank and Royal Bank of Scotland Â£5 and Â£10 polymer banknotes. Forensic Science International, 2020, 312, 110334.	1.3	7
7	Identifying blood-specific age-related DNA methylation markers on the Illumina MethylationEPICÂ® BeadChip. Forensic Science International, 2019, 303, 109944.	1.3	22
8	Estimating time since deposition using quantification of RNA degradation in body fluid-specific markers. Forensic Science International, 2019, 298, 58-63.	1.3	40
9	Evaluation of 30 insertion/deletion polymorphisms as forensic markers in the Kuwaiti population. Forensic Science International: Genetics Supplement Series, 2019, 7, 737-738.	0.1	1
10	A multi-tissue age prediction model based on DNA methylation analysis. Forensic Science International: Genetics Supplement Series, 2017, 6, e62-e64.	0.1	10
11	Quantification of RNA degradation of blood-specific markers to indicate the age of bloodstains. Forensic Science International: Genetics Supplement Series, 2017, 6, e453-e455.	0.1	14
12	Variation in the intensity of selection on codon bias over time causes contrasting patterns of base composition evolution in <i>Drosophila</i> . Genome Biology and Evolution, 2017, 9, eww291.	1.1	38
13	Strain-specific and pooled genome sequences for populations of <i>Drosophila melanogaster</i> from three continents.. F1000Research, 2015, 4, 31.	0.8	15
14	The Discovery, Distribution, and Evolution of Viruses Associated with <i>Drosophila melanogaster</i> . PLoS Biology, 2015, 13, e1002210.	2.6	272
15	The Relation between Recombination Rate and Patterns of Molecular Evolution and Variation in <i>Drosophila melanogaster</i> . Molecular Biology and Evolution, 2014, 31, 1010-1028.	3.5	144
16	Estimation of the Spontaneous Mutation Rate per Nucleotide Site in a <i>Drosophila melanogaster</i> Full-Sib Family. Genetics, 2014, 196, 313-320.	1.2	248
17	Codon Usage Bias and Effective Population Sizes on the X Chromosome versus the Autosomes in <i>Drosophila melanogaster</i> . Molecular Biology and Evolution, 2013, 30, 811-823.	3.5	41
18	Variation in male and female mating behaviour among different populations of the two-spot ladybird, <i>Adalia bipunctata</i> (Coleoptera: Coccinellidae). European Journal of Entomology, 2013, 110, 87-93.	1.2	10

#	ARTICLE	IF	CITATIONS
19	Molecular Evolution in Nonrecombining Regions of the <i>Drosophila melanogaster</i> Genome. <i>Genome Biology and Evolution</i> , 2012, 4, 278-288.	1.1	51
20	Ancestral polymorphisms in <i>Drosophila pseudoobscura</i> and <i>Drosophila miranda</i> . <i>Genetical Research</i> , 2011, 93, 255-263.	0.3	10
21	Determinants of Synonymous and Nonsynonymous Variability in Three Species of <i>Drosophila</i> . <i>Molecular Biology and Evolution</i> , 2011, 28, 1731-1743.	3.5	36
22	Genetic Diversity, Population Structure and Wolbachia Infection Status in a Worldwide Sample of <i>Drosophila melanogaster</i> and <i>D. simulans</i> Populations. <i>PLoS ONE</i> , 2011, 6, e26318.	1.1	44
23	Estimating the Parameters of Selection on Nonsynonymous Mutations in <i>Drosophila pseudoobscura</i> and <i>D. miranda</i> . <i>Genetics</i> , 2010, 185, 1381-1396.	1.2	61
24	Female multiple mating in wild and laboratory populations of the two-spot ladybird, <i>Adalia bipunctata</i> . <i>Molecular Ecology</i> , 2008, 17, 3189-3197.	2.0	39
25	Non-neutral processes drive the nucleotide composition of non-coding sequences in <i>Drosophila</i> . <i>Biology Letters</i> , 2008, 4, 438-441.	1.0	40
26	Elevated levels of expression associated with regions of the <i>Drosophila</i> genome that lack crossing over. <i>Biology Letters</i> , 2008, 4, 758-761.	1.0	15
27	Positive and Negative Selection on Noncoding DNA in <i>Drosophila simulans</i> . <i>Molecular Biology and Evolution</i> , 2008, 25, 1825-1834.	3.5	91
28	A multispecies approach for comparing sequence evolution of X-linked and autosomal sites in <i>Drosophila</i> . <i>Genetical Research</i> , 2008, 90, 421-431.	0.3	29
29	The age and evolution of an antiviral resistance mutation in <i>Drosophila melanogaster</i> . <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2007, 274, 2027-2034.	1.2	48
30	Reduced efficacy of selection in regions of the <i>Drosophila</i> genome that lack crossing over. <i>Genome Biology</i> , 2007, 8, R18.	13.9	140
31	Temporal effects of multiple mating on components of fitness in the two-spot ladybird, <i>Adalia bipunctata</i> (Coleoptera: Coccinellidae). <i>European Journal of Entomology</i> , 2007, 104, 393-398.	1.2	23
32	Multilocus patterns of nucleotide variability and the demographic and selection history of <i>Drosophila melanogaster</i> populations. <i>Genome Research</i> , 2005, 15, 790-799.	2.4	247
33	Patterns of intron sequence evolution in <i>Drosophila</i> are dependent upon length and GC content. <i>Genome Biology</i> , 2005, 6, R67.	13.9	158