

Tomasz Kupiec

List of Publications by Year in descending order

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Version: 2024-02-01

38
papers

1,422
citations

516710

16
h-index

330143

37
g-index

41
all docs

41
docs citations

41
times ranked

1681
citing authors

#	ARTICLE	IF	CITATIONS
1	Confirmation of Paternity despite Three Genetic Incompatibilities at Chromosome 2. <i>Genes</i> , 2021, 12, 62.	2.4	3
2	Recommendations of the Polish Speaking Working Group of the International Society for Forensic Genetics on forensic Y chromosome typing. <i>Archiwum Medycyny Sadowej I Kryminologii</i> , 2020, 70, 1-18.	0.3	2
3	Evaluation of the performance of the beta version of the ForenSeq DNA signature Prep Kit on the MiSeq FGx forensic genomics system. <i>Forensic Science International: Genetics Supplement Series</i> , 2019, 7, 585-586.	0.3	3
4	Y chromosome sequence variation of common forensic STR markers and their flanking regions among Polish population. <i>Forensic Science International: Genetics Supplement Series</i> , 2019, 7, 557-560.	0.3	1
5	Dual amplification strategy for improved efficiency of forensic DNA analysis using NGM Detectâ„¢, NGMâ„¢ or Globalfilerâ„¢ kits. <i>Forensic Science International: Genetics</i> , 2018, 35, 46-49.	3.1	8
6	Recommendations of the Polish Speaking Working Group of the International Society for Forensic Genetics for forensic mitochondrial DNA testing. <i>Archiwum Medycyny Sadowej I Kryminologii</i> , 2018, 68, 242-258.	0.3	1
7	A case study of an unknown mass grave " Hostages killed 70 years ago by a Nazi firing squad identified thanks to genetics. <i>Forensic Science International</i> , 2017, 278, 173-176.	2.2	12
8	Genetic Identification of Communist Crimes™ Victims (1944–1956) Based on the Analysis of One of Many Mass Graves Discovered on the Powazki Military Cemetery in Warsaw, Poland. <i>Journal of Forensic Sciences</i> , 2016, 61, 1450-1455.	1.6	14
9	Simultaneous Whole Mitochondrial Genome Sequencing with Short Overlapping Amplicons Suitable for Degraded DNA Using the Ion Torrent Personal Genome Machine. <i>Human Mutation</i> , 2015, 36, 1236-1247.	2.5	51
10	Development of a forensically useful age prediction method based on DNA methylation analysis. <i>Forensic Science International: Genetics</i> , 2015, 17, 173-179.	3.1	236
11	Practical aspects of genetic identification of hallucinogenic and other poisonous mushrooms for clinical and forensic purposes. <i>Croatian Medical Journal</i> , 2015, 56, 32-40.	0.7	7
12	Examination of DNA methylation status of the ELOVL2 marker may be useful for human age prediction in forensic science. <i>Forensic Science International: Genetics</i> , 2015, 14, 161-167.	3.1	163
13	A new dimension of the forensic DNA expertise " the need for training experts and expertise recipients. <i>Archiwum Medycyny Sadowej I Kryminologii</i> , 2014, 3, 175-194.	0.3	4
14	Collaborative EDNAP exercise on the IrisPlex system for DNA-based prediction of human eye colour. <i>Forensic Science International: Genetics</i> , 2014, 11, 241-251.	3.1	23
15	Bona fide colour: DNA prediction of human eye and hair colour from ancient and contemporary skeletal remains. <i>Investigative Genetics</i> , 2013, 4, 3.	3.3	58
16	A cautionary note on switching mitochondrial DNA reference sequences in forensic genetics. <i>Forensic Science International: Genetics</i> , 2012, 6, e182-e184.	3.1	24
17	Prediction of Eye Color from Genetic Data Using Bayesian Approach*. <i>Journal of Forensic Sciences</i> , 2012, 57, 880-886.	1.6	30
18	Gene–gene interactions contribute to eye colour variation in humans. <i>Journal of Human Genetics</i> , 2011, 56, 447-455.	2.3	57

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19	Model-based prediction of human hair color using DNA variants. <i>Human Genetics</i> , 2011, 129, 443-454.	3.8	151
20	Genetic examination of the putative skull of Jan Kochanowski reveals its female sex. <i>Croatian Medical Journal</i> , 2011, 52, 403-409.	0.7	5
21	Genetic variation of 15 autosomal STR loci in a population sample from Poland. <i>Legal Medicine</i> , 2010, 12, 246-248.	1.3	16
22	Genetic identification of putative remains of the famous astronomer Nicolaus Copernicus. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 12279-12282.	7.1	46
23	A population data for 17 Y-chromosome STR loci in South Poland population sample—Some DYS458.2 variants uncovered and sequenced. <i>Forensic Science International: Genetics</i> , 2009, 4, e43-e44.	3.1	16
24	Association of the SLC45A2 gene with physiological human hair colour variation. <i>Journal of Human Genetics</i> , 2008, 53, 966-971.	2.3	60
25	Association of Polymorphic Sites in the <i>OCA2</i> Gene with Eye Colour Using the Tree Scanning Method. <i>Annals of Human Genetics</i> , 2008, 72, 184-192.	0.8	29
26	Examples of combining genetic evidence—Bayesian network approach. <i>Forensic Science International: Genetics Supplement Series</i> , 2008, 1, 669-670.	0.3	1
27	The <i>OCA2</i> gene as a marker for eye colour prediction. <i>Forensic Science International: Genetics Supplement Series</i> , 2008, 1, 536-537.	0.3	2
28	Application of BioRobot M48 to forensic DNA extraction. <i>Forensic Science International: Genetics Supplement Series</i> , 2008, 1, 58-59.	0.3	3
29	Determination of Phenotype Associated SNPs in the MC1R Gene. <i>Journal of Forensic Sciences</i> , 2007, 52, 349-354.	1.6	64
30	A search for genes modulated by interleukin-6 alone or with interleukin-1 β in HepG2 cells using differential display analysis. <i>Biochimica Et Biophysica Acta - Molecular Basis of Disease</i> , 2006, 1762, 319-328.	3.8	15
31	Determination of forensically relevant SNPs in the MC1R gene. <i>International Congress Series</i> , 2006, 1288, 816-818.	0.2	1
32	Distribution of mtDNA Haplogroups in a Population Sample from Poland. <i>Journal of Forensic Sciences</i> , 2005, 50, 1-2.	1.6	5
33	The EDNAP mitochondrial DNA population database (EMPOP) collaborative exercises: organisation, results and perspectives. <i>Forensic Science International</i> , 2004, 139, 215-226.	2.2	105
34	Beyond HV1 and HV2—identification of valuable mitochondrial DNA single nucleotide polymorphisms. <i>International Congress Series</i> , 2004, 1261, 100-102.	0.2	0
35	Validation of <i>Cytochrome b</i> Sequence Analysis as a Method of Species Identification. <i>Journal of Forensic Sciences</i> , 2003, 48, 1-5.	1.6	101
36	Homogeneity and distinctiveness of Polish paternal lineages revealed by Y chromosome microsatellite haplotype analysis. <i>Human Genetics</i> , 2002, 110, 592-600.	3.8	91

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37	STR data for SGM Plus and penta E and D loci in a population sample from south Poland. Forensic Science International, 2002, 127, 237-239.	2.2	8
38	STR data for AmpF/STR Profiler Plus loci in south Poland. Forensic Science International, 2001, 122, 173-174.	2.2	3