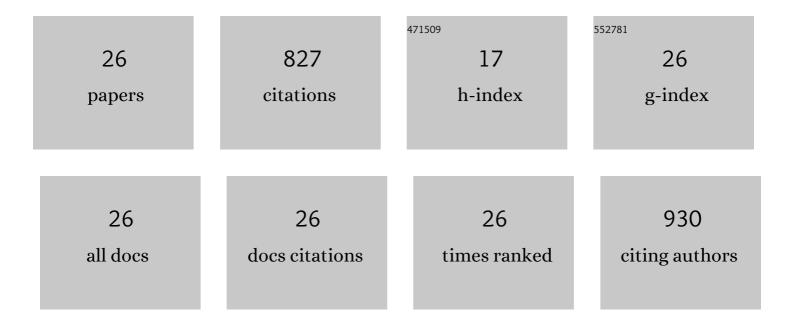
## Marcin WoÅ<sup>o</sup>niak

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Clear phylogeographic pattern and genetic structure of wild boar Sus scrofa population in Central and Eastern Europe. Scientific Reports, 2021, 11, 9680.	3.3	11
2	Recommendations of the Polish Speaking Working Group of the International Society for Forensic Genetics on forensic Y chromosome typing. Archiwum Medycyny Sadowej I Kryminologii, 2020, 70, 1-18.	0.3	2
3	Raoultella spp. – reliable identification, susceptibility to antimicrobials and antibiotic resistance mechanisms. Journal of Medical Microbiology, 2020, 69, 233-238.	1.8	11
4	Intra- and inter-population analysis of haplotype diversity in Yfiler ® Plus system using a wide set of representative data from Polish population. Forensic Science International: Genetics, 2017, 28, e22-e25.	3.1	20
5	Y chromosome haplotype diversity in Mongolic-speaking populations and gene conversion at the duplicated STR DYS385a,b in haplogroup C3-M407. Journal of Human Genetics, 2016, 61, 491-496.	2.3	7
6	A novel multiplex assay amplifying 13 Y-STRs characterized by rapid and moderate mutation rate. Forensic Science International: Genetics, 2015, 15, 49-55.	3.1	19
7	Genetic data from Y chromosome STR and SNP loci in Ukrainian population. Forensic Science International: Genetics, 2013, 7, 200-203.	3.1	25
8	Y-chromosome diversity in the Kalmyks at the ethnical and tribal levels. Journal of Human Genetics, 2013, 58, 804-811.	2.3	18
9	Y-chromosome variation in Tajiks and Iranians. Annals of Human Biology, 2013, 40, 48-54.	1.0	6
10	The Y-chromosome C3* Star-Cluster Attributed to Genghis Khan's Descendants is Present at High Frequency in the Kerey Clan from Kazakhstan. Human Biology, 2012, 84, 79-89.	0.2	39
11	Ancient links between Siberians and Native Americans revealed by subtyping the Y chromosome haplogroup Q1a. Journal of Human Genetics, 2011, 56, 583-588.	2.3	56
12	Similarities and distinctions in Y chromosome gene pool of Western Slavs. American Journal of Physical Anthropology, 2010, 142, 540-548.	2.1	27
13	Phylogeography of the Yâ€chromosome haplogroup C in northern Eurasia. Annals of Human Genetics, 2010, 74, 539-546.	0.8	45
14	Developing STR databases on structured populations: The native South Siberian population versus the Russian population. Forensic Science International: Genetics, 2009, 3, e111-e116.	3.1	22
15	Analysis of forensically used autosomal short tandem repeat markers in Polish and neighboring populations. Forensic Science International: Genetics, 2008, 2, 205-211.	3.1	21
16	Continuity of Y chromosome haplotypes in the population of Southern Poland before and after the Second World War. Forensic Science International: Genetics, 2007, 1, 134-140.	3.1	22
17	Complex interactions of the Eastern and Western Slavic populations with other European groups as revealed by mitochondrial DNA analysis. Forensic Science International: Genetics, 2007, 1, 141-147.	3.1	60
18	Y-chromosome haplogroup N dispersals from south Siberia to Europe. Journal of Human Genetics, 2007, 52, 763-770.	2.3	65

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#	Article	IF	CITATIONS
19	Allelic and haplotypic frequencies at 11 Y-STR loci in Buryats from South-East Siberia. Forensic Science International, 2006, 164, 271-275.	2.2	20
20	Contrasting patterns of Y-chromosome variation in South Siberian populations from Baikal and Altai-Sayan regions. Human Genetics, 2006, 118, 591-604.	3.8	70
21	Significant genetic differentiation between Poland and Germany follows present-day political borders, as revealed by Y-chromosome analysis. Human Genetics, 2005, 117, 428-443.	3.8	123
22	Homogeneity and distinctiveness of Polish paternal lineages revealed by Y chromosome microsatellite haplotype analysis. Human Genetics, 2002, 110, 592-600.	3.8	91
23	Population genetics of the STRs vWA, TH01, TPOX, CSF1PO, D5S818, D13S317, D7S820, D16S539, LPL, F13B, FESFPS, F13A01 and ACTBP2 in the Pomerania-Kujawy region of Poland. Forensic Science International, 2001, 119, 119-122.	2.2	11
24	Screening of a highly polymorphic microsatellite for microheterogeneity in human identification. Electrophoresis, 1998, 19, 667-670.	2.4	2
25	Novel BRCA1 mutations and more frequent intron-20 alteration found among 236 women from Western Poland. Oncogene, 1997, 15, 1773-1779.	5.9	28
26	Optimization of a hexaplex DNA amplification from short tandem repeat and amelogenin loci. Electrophoresis, 1997, 18, 1627-1632.	2.4	6