Oliver V StojkoviÄ

List of Publications by Year
in descending order

33	1,294	759233 12	454955 30
papers	citations	h-index	g-index

35	35	35	2395
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Dominant-negative mutations in the DNA-binding domain of STAT3 cause hyper-IgE syndrome. Nature, 2007, 448, 1058-1062.	27.8	930
2	Allele frequencies and population data for 17 Y-chromosome STR loci in a Serbian population sample from Vojvodina province. Forensic Science International, 2008, 176, e23-e28.	2.2	31
3	250 CTG repeats in DMPK is a threshold for correlation of expansion size and age at onset of juvenile-adult DM1. Human Mutation, 2002, 19, 131-139.	2.5	30
4	Human Yâ€chromosome short tandem repeats: A tale of acculturation and migrations as mechanisms for the diffusion of agriculture in the Balkan Peninsula. American Journal of Physical Anthropology, 2010, 142, 380-390.	2.1	29
5	DNA typing from handled items. Forensic Science International: Genetics Supplement Series, 2008, 1, 411-412.	0.3	26
6	Y-STR profiling in two Afghanistan populations. Legal Medicine, 2011, 13, 103-108.	1.3	26
7	Prediction of autosomal STR typing success in ancient and Second World War bone samples. Forensic Science International: Genetics, 2017, 27, 17-26.	3.1	26
8	Laboratory Evolution of Life-History Traits in the Bean Weevil (Acanthoscelides obtectus): The Effects of Density-Dependent and Age- Specific Selection. Evolution; International Journal of Organic Evolution, 1997, 51, 1896.	2.3	24
9	Y-chromosomal microsatellite diversity in three culturally defined regions of historical Tibet. Forensic Science International: Genetics, 2012, 6, 437-446.	3.1	20
10	Divergent patrilineal signals in three Roma populations. American Journal of Physical Anthropology, 2011, 144, 80-91.	2.1	16
11	Y-STR diversity in the Himalayas. International Journal of Legal Medicine, 2011, 125, 367-375.	2.2	15
12	Correlation between triplet repeat expansion and computed tomography measures of caudate nuclei atrophy in Huntington's disease. Journal of Neurology, 1999, 246, 1090-1093.	3.6	13
13	Polymorphisms in<i> ACE</i> and <i>ACTN3</i> Genes and Blood Pressure Response to Acute Exercise in Elite Male Athletes from Serbia. Tohoku Journal of Experimental Medicine, 2017, 243, 311-320.	1.2	13
14	Case Report: Post-mortem Histopathological and Molecular Analyses of the Very First Documented COVID-19-Related Death in Europe. Frontiers in Medicine, 2021, 8, 612758.	2.6	12
15	Forensic DNA databases in Western Balkan region: retrospectives, perspectives, and initiatives. Croatian Medical Journal, 2011, 52, 235-244.	0.7	11
16	Genetics of suspected thrombophilia in Serbian females with infertility, including three cases, homozygous for<i>FII</i>20210A or<i>FV</i>1691A mutations. Human Fertility, 2017, 20, 132-139.	1.7	11
17	Polymorphisms and haplotypes in VDR gene are associated with female idiopathic infertility. Human Fertility, 2020, 23, 101-110.	1.7	11
18	The Status of SCA1, MJD/SCA3, FRDA, DRPLA and MD Triplet Containing Genes in Patients with Huntington Disease and Healthy Controls. Journal of Neurogenetics, 2000, 14, 257-263.	1.4	9

#	Article	IF	Citations
19	Defining screening panel of functional variants of CYP1A1, CYP2C9, CYP2C19, CYP2D6, and CYP3A4 genes in Serbian population. International Journal of Legal Medicine, 2020, 134, 433-439.	2.2	7
20	Is the 31 CAG repeat allele of the spinocerebellar ataxia 1 (SCA1) gene locus non-specifically associated with trinucleotide expansion diseases?. Psychiatric Genetics, 2001, 11, 201-205.	1.1	5
21	Multidisciplinary investigation links backward-speech trait and working memory through genetic mutation. Scientific Reports, 2016, 6, 20369.	3.3	5
22	Cryptic Diversity of the European Blind Mole Rat Nannospalax leucodon Species Complex: Implications for Conservation. Animals, 2022, 12, 1097.	2.3	5
23	On the Bantu expansion. Gene, 2016, 593, 48-57.	2.2	4
24	Yugoslav population data on nine STR loci. Forensic Science International, 2001, 115, 239-240.	2.2	3
25	16S rRNA gene polymorphism supports cryptic speciation within the lesser blind mole rat Nannospalax leucodon superspecies (Rodentia: Spalacidae). Mammalian Biology, 2020, 100, 315-324.	1.5	3
26	Poly(A) tailing of ancient DNA: a method for reproducible microsatellite genotyping. Analytical Biochemistry, 2003, 318, 124-131.	2.4	2
27	Population Data on HLA-DQA1, LDLR, GYPA, HBGG, D7S8, and GC PCR-Based Loci in Serbia. Journal of Forensic Sciences, 2006, 51, 699-699.	1.6	2
28	Combinations of fibrinolytic gene polymorphisms (plasminogen activator inhibitor type 1 4G/5G, factor) Tj ETQq0 0 0 rgBT /Overlock 10 Coagulation and Fibrinolysis, 2021, 32, 103-107.	1.0	2
29	Hemostasis-related gene polymorphisms and their epistatic relationship in women with idiopathic infertility. Blood Coagulation and Fibrinolysis, 2019, 30, 253-262.	1.0	1
30	Monitoring of Pediatric Patients With Malignant Hematological Diseases After Allogeneic HSCT. Journal of Pediatric Hematology/Oncology, 2012, 34, e253-e257.	0.6	0
31	STR loci D19S216, D20S502 and D20S842 analysis in the Serbian population using dentin DNA. Archives of Biological Sciences, 2011, 63, 55-58.	0.5	0
32	The angiotensin converting enzyme (ace) gene polymorphism: Insight study of the renal regulation of the arterial blood pressure. Medicinski Podmladak, 2016, 67, 31-35.	0.0	0
33	Should MTHFR 1298 A>C be tested together with MTHFR 677 C>T polymorphism in women with reproductive challenges?. Genetika, 2017, 49, 377-386.	0.4	0