Bram Bekaert

List of Publications by Year in descending order

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566801 454577 37 968 15 30 citations h-index g-index papers 39 39 39 1461 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Improved age determination of blood and teeth samples using a selected set of DNA methylation markers. Epigenetics, 2015, 10, 922-930.	1.3	187
2	Maternal intake of methyl-group donors affects DNA methylation of metabolic genes in infants. Clinical Epigenetics, 2017, 9, 16.	1.8	129
3	Dietary and supplemental maternal methyl-group donor intake and cord blood DNA methylation. Epigenetics, 2017, 12, 1-10.	1.3	112
4	Randomized controlled trial of the effect of selenium supplementation on thyroid function in the elderly in the United Kingdom. American Journal of Clinical Nutrition, 2008, 87, 370-378.	2.2	97
5	Changes in DNA methylation induced by multi-walled carbon nanotube exposure in the workplace. Nanotoxicology, 2017, 11, 1195-1210.	1.6	41
6	Evaluation of three statistical prediction models for forensic age prediction based on DNA methylation. Forensic Science International: Genetics, 2018, 34, 128-133.	1.6	37
7	The Influence of the Duration of Breastfeeding on the Infant's Metabolic Epigenome. Nutrients, 2019, 11, 1408.	1.7	29
8	Carbon Nanotube- and Asbestos-Induced DNA and RNA Methylation Changes in Bronchial Epithelial Cells. Chemical Research in Toxicology, 2019, 32, 850-860.	1.7	28
9	DNA Methylation and Brainâ€Derived Neurotrophic Factor Expression Account for Symptoms and Widespread Hyperalgesia in Patients With Chronic Fatigue Syndrome and Comorbid Fibromyalgia. Arthritis and Rheumatology, 2020, 72, 1936-1944.	2.9	28
10	Increased methylation of NR3C1 and SLC6A4 is associated with blunted cortisol reactivity to stress in major depression. Neurobiology of Stress, 2020, 13, 100272.	1.9	25
11	Forensic Epigenetic Age Estimation and Beyond: Ethical and Legal Considerations. Trends in Genetics, 2018, 34, 489-491.	2.9	24
12	A selective set of DNA-methylation markers for age determination of blood, teeth and buccal samples. Forensic Science International: Genetics Supplement Series, 2015, 5, e144-e145.	0.1	22
13	Effect of selenium status and supplementation with highâ€selenium yeast on plasma homocysteine and B vitamin concentrations in the UK elderly. Molecular Nutrition and Food Research, 2008, 52, 1324-1333.	1.5	21
14	The effect of paternal methyl-group donor intake on offspring DNA methylation and birth weight. Journal of Developmental Origins of Health and Disease, 2017, 8, 311-321.	0.7	21
15	Single-walled and multi-walled carbon nanotubes induce sequence-specific epigenetic alterations in 16 HBE cells. Oncotarget, 2018, 9, 20351-20365.	0.8	21
16	Allele frequencies for the new European Standard Set (ESS) loci and D1S1677 in the Belgian population. Forensic Science International: Genetics, 2012, 6, e75-e77.	1.6	16
17	Epigenetic perspective on the role of brain-derived neurotrophic factor in burnout. Translational Psychiatry, 2020, 10, 354.	2.4	15
18	Development and evaluation of multiplex Y-STR assays for application in molecular genealogy. Forensic Science International: Genetics Supplement Series, 2009, 2, 57-59.	0.1	12

#	Article	IF	CITATIONS
19	Automated DNA extraction of single dog hairs without roots for mitochondrial DNA analysis. Forensic Science International: Genetics, 2012, 6, 277-281.	1.6	12
20	Biohistorical materials and contemporary privacy concerns-the forensic case of King Albert I. Forensic Science International: Genetics, 2016, 24, 202-210.	1.6	11
21	New aspects of dental implants and DNA technology in human identification. Forensic Science International, 2019, 302, 109926.	1.3	11
22	Survival of forensic trace evidence on improvised explosive devices: perspectives on individualisation. Scientific Reports, 2020, 10, 12813.	1.6	11
23	Automating a combined composite–consensus method to generate DNA profiles from low and high template mixture samples. Forensic Science International: Genetics, 2012, 6, 588-593.	1.6	10
24	Plasma selenium concentration and prostate cancer risk. American Journal of Clinical Nutrition, 2009, 89, 1276-1277.	2.2	8
25	Role of NR3C1 and SLC6A4 methylation in the HPA axis regulation in burnout. Journal of Affective Disorders, 2021, 295, 505-512.	2.0	7
26	Increased sensitivity for amplified STR alleles on capillary sequencers with BigDye® XTerminatorâ,,¢. Forensic Science International: Genetics Supplement Series, 2009, 2, 123-124.	0.1	6
27	A comparison of mtDNA and Y chromosome diversity in Malay populations. International Congress Series, 2006, 1288, 252-255.	0.2	4
28	Validation of a microchip electrophoresis system as a DNA amplification control. Forensic Science International: Genetics Supplement Series, 2009, 2, 119-120.	0.1	3
29	Multiplex DNA amplification and barcoding in a single reaction for 454 Roche sequencing: A comprehensive study on the control region of the mitochondrial genome. Forensic Science International: Genetics Supplement Series, 2013, 4, e111-e112.	0.1	3
30	The AMOVA analyses and phylogenetic relationships of Pakistani population using Y STRs. International Congress Series, 2006, 1288, 171-173.	0.2	2
31	An automated approach for generating consensus profiles from low template STR typing results. Forensic Science International: Genetics Supplement Series, 2011, 3, e435-e436.	0.1	2
32	The development of a forensic clock to determine time of death. Forensic Science International: Genetics Supplement Series, 2017, 6, e162-e163.	0.1	2
33	Cell survival and DNA damage repair are promoted in the human blood thanatotranscriptome shortly after death. Scientific Reports, 2021, 11, 16585.	1.6	2
34	Evaluation of infrared photography for latent bloodstain visualization and the influence of time. Forensic Science International, 2022, 331, 111167.	1.3	2
35	Influence of ink and smoke ATM security systems on dactyloscopy and subsequent DNA analysis after detonation. Forensic Science International: Genetics, 2021, 54, 102540.	1.6	1
36	Optimization and validation of the SNPforID 34-SNPplex for POP7â,,¢. Forensic Science International: Genetics Supplement Series, 2011, 3, e43-e44.	0.1	0

ARTICLE IF CITATIONS

37 O18-1â€...Epigenetic effects of occupational exposure to carbon nanotubes., 2016,,. o