Thore Egeland

List of Publications by Year in descending order

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THORE FORME

#	Article	IF	CITATIONS
1	Pairwise relatedness testing in the context of inbreeding: expectation and variance of the likelihood ratio. International Journal of Legal Medicine, 2021, 135, 117-129.	1.2	5
2	Joint DNA-based disaster victim identification. Scientific Reports, 2021, 11, 13661.	1.6	4
3	Strategies for pairwise searches in forensic kinship analysis. Forensic Science International: Genetics, 2021, 54, 102562.	1.6	5
4	Making decisions in missing person identification cases with low statistical power. Forensic Science International: Genetics, 2021, 54, 102519.	1.6	6
5	Prioritising family members for genotyping in missing person cases: A general approach combining the statistical power of exclusion and inclusion. Forensic Science International: Genetics, 2020, 49, 102376.	1.6	16
6	The failing measurement of attitudes: How semantic determinants of individual survey responses come to replace measures of attitude strength. Behavior Research Methods, 2018, 50, 2345-2365.	2.3	21
7	Degradation in forensic trace DNA samples explored by massively parallel sequencing. Forensic Science International: Genetics, 2017, 27, 160-166.	1.6	16
8	DNA Commission of the International Society for Forensic Genetics (ISFG): Guidelines on the use of X-STRs in kinship analysis. Forensic Science International: Genetics, 2017, 29, 269-275.	1.6	71
9	Exact likelihood ratio calculations for pairwise cases. Forensic Science International: Genetics, 2017, 29, 218-224.	1.6	3
10	Key individuals for discerning pedigrees belonging to the same autosomal kinship class. Forensic Science International: Genetics, 2017, 29, 71-79.	1.6	4
11	The implications of shedder status and background DNA on direct and secondary transfer in an attack scenario. Forensic Science International: Genetics, 2017, 29, 48-60.	1.6	80
12	Characterization of degradation and heterozygote balance by simulation of the forensic DNA analysis process. International Journal of Legal Medicine, 2017, 131, 303-317.	1.2	25
13	Evaluating the statistical power of DNA-based identification, exemplified by †The missing grandchildren of Argentina'. Forensic Science International: Genetics, 2017, 31, 57-66.	1.6	17
14	Searching for relationships. , 2016, , 51-84.		0
15	Dependent markers. , 2016, , 85-129.		Ο
16	Mixtures with relatives and linked markers. International Journal of Legal Medicine, 2016, 130, 621-634.	1.2	9
17	Contamination during criminal investigation: Detecting police contamination and secondary DNA transfer from evidence bags. Forensic Science International: Genetics, 2016, 23, 121-129.	1.6	37
18	About the number of contributors to a forensic sample. Forensic Science International: Genetics, 2016, 25, e18-e19.	1.6	11

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19	The likelihood ratio as a random variable for linked markers in kinship analysis. International Journal of Legal Medicine, 2016, 130, 1445-1456.	1.2	5
20	Kinship. Security Science and Technology, 2016, , 81-100.	0.5	0
21	Knowing the midwife before delivery reduces the prevalence of caesarean section on demand in a group of second time mothers with a complicated first delivery. Nordic Journal of Nursing Research, 2016, 36, 44-50.	0.6	1
22	Lipoprotein subfractions by nuclear magnetic resonance are associated with tumor characteristics in breast cancer. Lipids in Health and Disease, 2016, 15, 56.	1.2	37
23	Relationship inference based on DNA mixtures. International Journal of Legal Medicine, 2016, 130, 323-329.	1.2	16
24	Exclusion probabilities and likelihood ratios with applications to mixtures. International Journal of Legal Medicine, 2016, 130, 39-57.	1.2	19
25	Models and implementation for relationship problems with dropout. International Journal of Legal Medicine, 2015, 129, 411-423.	1.2	9
26	Secondary and subsequent DNA transfer during criminal investigation. Forensic Science International: Genetics, 2015, 17, 155-162.	1.6	75
27	Genotyping and interpretation of STR-DNA: Low-template, mixtures and database matches—Twenty years of research and development. Forensic Science International: Genetics, 2015, 18, 100-117.	1.6	116
28	High-Density Lipoprotein-Cholesterol, Daily Estradiol and Progesterone, and Mammographic Density Phenotypes in Premenopausal Women. Cancer Prevention Research, 2015, 8, 535-544.	0.7	10
29	Gene variations in oestrogen pathways, CYP19A1, daily 17β-estradiol and mammographic density phenotypes in premenopausal women. Breast Cancer Research, 2014, 16, 499.	2.2	12
30	A general approach to power calculation for relationship testing. Forensic Science International: Genetics, 2014, 9, 186-190.	1.6	30
31	Mixtures with relatives: A pedigree perspective. Forensic Science International: Genetics, 2014, 10, 49-54.	1.6	14
32	STR-validator: An open source platform for validation and process control. Forensic Science International: Genetics, 2014, 13, 154-166.	1.6	16
33	Familias 3 – Extensions and new functionality. Forensic Science International: Genetics, 2014, 13, 121-127.	1.6	156
34	Exact computation of the distribution of likelihood ratios with forensic applications. Forensic Science International: Genetics, 2014, 9, 93-101.	1.6	31
35	Regression models for DNA-mixtures. Forensic Science International: Genetics, 2014, 11, 105-110.	1.6	3
36	On the meaning of the likelihood ratio: Is a large number always an indication of strength of evidence?. Forensic Science International: Genetics Supplement Series, 2013, 4, e176-e177.	0.1	5

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37	FamLink – A user friendly software for linkage calculations in family genetics. Forensic Science International: Genetics, 2012, 6, 616-620.	1.6	41
38	A Statistical Framework for the Interpretation of mtDNA Mixtures: Forensic and Medical Applications. PLoS ONE, 2011, 6, e26723.	1.1	11
39	A CLEC16A variant confers risk for juvenile idiopathic arthritis and anti-cyclic citrullinated peptide antibody negative rheumatoid arthritis. Annals of the Rheumatic Diseases, 2010, 69, 1471-1474.	0.5	55
40	Mutation screening of PTPN22: association of the 1858T-allele with Addison's disease. European Journal of Human Genetics, 2008, 16, 977-982.	1.4	81
41	Polymorphisms in <i>CLEC16A</i> and <i>ClITA</i> at 16p13 Are Associated with Primary Adrenal Insufficiency. Journal of Clinical Endocrinology and Metabolism, 2008, 93, 3310-3317.	1.8	108
42	Estimating Haplotype Frequency and Coverage of Databases. PLoS ONE, 2008, 3, e3988.	1.1	29
43	The DNA Database Search Controversy Revisited: Bridging the Bayesian-Frequentist Gap. Biometrics, 2007, 63, 922-925.	0.8	24
44	Insulin Resistance after Renal Transplantation: The Effect of Steroid Dose Reduction and Withdrawal. Journal of the American Society of Nephrology: JASN, 2004, 15, 3233-3239.	3.0	158
45	Breech birth at term: vaginal delivery or elective cesarean section? A systematic review of the literature by a Norwegian review team. Acta Obstetricia Et Gynecologica Scandinavica, 2004, 83, 126-130.	1.3	21
46	Determinants of insulin secretion after renal transplantation. Metabolism: Clinical and Experimental, 2003, 52, 573-578.	1.5	46
47	Response to Montori et al Diabetes Care, 2002, 25, 1667-1667.	4.3	1
48	The impact of impaired insulin release and insulin resistance on glucose intolerance after renal transplantation*. Clinical Transplantation, 2002, 16, 389-396.	0.8	59
49	Tapering off prednisolone and cyclosporin the first year after renal transplantation: the effect on glucose tolerance. Nephrology Dialysis Transplantation, 2001, 16, 829-835.	0.4	121
50	Metabolic cardiovascular syndrome after renal transplantation. Nephrology Dialysis Transplantation, 2001, 16, 1047-1052.	0.4	63
51	GLUCOSE INTOLERANCE AFTER RENAL TRANSPLANTATION DEPENDS UPON PREDNISOLONE DOSE AND RECIPIENT AGE1. Transplantation, 1997, 64, 979-983.	0.5	273