Brahim Aissani

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28 871 14 28 g-index

28 g-index

28 ext. papers ext. citations avg, IF L-index

#	Paper	IF	Citations
28	The distribution of genes in the human genome. <i>Gene</i> , 1991 , 100, 181-7	3.8	223
27	Correlations between the compositional properties of human genes, codon usage, and amino acid composition of proteins. <i>Journal of Molecular Evolution</i> , 1991 , 32, 504-10	3.1	156
26	CpG islands: features and distribution in the genomes of vertebrates. <i>Gene</i> , 1991 , 106, 173-83	3.8	91
25	CpG islands, genes and isochores in the genomes of vertebrates. <i>Gene</i> , 1991 , 106, 185-95	3.8	80
24	The compositional properties of human genes. <i>Journal of Molecular Evolution</i> , 1991 , 32, 493-503	3.1	61
23	Host genetics and HIV-1 viral load set-point in African-Americans. <i>Aids</i> , 2009 , 23, 673-7	3.5	31
22	Compositional mapping of the human dystrophin-encoding gene. <i>Gene</i> , 1992 , 122, 329-35	3.8	28
21	A genome-wide association study of host genetic determinants of the antibody response to Anthrax Vaccine Adsorbed. <i>Vaccine</i> , 2012 , 30, 4778-84	4.1	21
20	A quantitative trait locus for body fat on chromosome 1q43 in French Canadians: linkage and association studies. <i>Obesity</i> , 2006 , 14, 1605-15	8	20
19	The major histocompatibility complex conserved extended haplotype 8.1 in AIDS-related non-Hodgkin lymphoma. <i>Journal of Acquired Immune Deficiency Syndromes (1999)</i> , 2009 , 52, 170-9	3.1	19
18	Follow-up to genome-wide linkage and admixture mapping studies implicates components of the extracellular matrix in susceptibility to and size of uterine fibroids. <i>Fertility and Sterility</i> , 2015 , 103, 528	-34.e1	3 ¹⁷
17	Evaluation of GWAS candidate susceptibility loci for uterine leiomyoma in the multi-ethnic NIEHS uterine fibroid study. <i>Frontiers in Genetics</i> , 2015 , 6, 241	4.5	15
16	Interleukin-10 (IL-10) pathway: genetic variants and outcomes of HIV-1 infection in African American adolescents. <i>PLoS ONE</i> , 2010 , 5, e13384	3.7	14
15	Multiple hits for the association of uterine fibroids on human chromosome 1q43. <i>PLoS ONE</i> , 2013 , 8, e58399	3.7	14
14	Mitochondrial DNA diversity in the African American population. <i>Mitochondrial DNA</i> , 2015 , 26, 445-51		13
13	Confounding by linkage disequilibrium. <i>Journal of Human Genetics</i> , 2014 , 59, 110-5	4.3	12
12	Compositional properties of telomeric regions from human chromosomes. FEBS Letters, 1991, 295, 22-	6 3.8	11

LIST OF PUBLICATIONS

11	Admixture mapping of genetic variants for uterine fibroids. <i>Journal of Human Genetics</i> , 2015 , 60, 533-8	4.3	10
10	Structure of tumor necrosis factor-alpha haploblocks in European populations. <i>Immunogenetics</i> , 2013 , 65, 543-52	3.2	9
9	Genetic determinants of uterine fibroid size in the multiethnic NIEHS uterine fibroid study. <i>International Journal of Molecular Epidemiology and Genetics</i> , 2015 , 6, 9-19	0.9	9
8	The utility of mitochondrial and y chromosome phylogenetic data to improve correction for population stratification. <i>Frontiers in Genetics</i> , 2012 , 3, 301	4.5	6
7	A candidate gene approach for virally induced cancer with application to HIV-related Kaposias sarcoma. <i>International Journal of Cancer</i> , 2014 , 134, 397-404	7.5	5
6	Fine mapping of the uterine leiomyoma locus on 1q43 close to a lncRNA in the RGS7-FH interval. <i>Endocrine-Related Cancer</i> , 2015 , 22, 633-43	5.7	2
5	Fine Mapping of the Body Fat QTL on Human Chromosome 1q43. <i>PLoS ONE</i> , 2016 , 11, e0153794	3.7	2
4	Mitochondrial DNA variation and virologic and immunological HIV outcomes in African Americans. <i>Aids</i> , 2014 , 28, 1871-8	3.5	1
3	Increasing Levels of Serum Heat Shock Protein 70 Precede the Development of AIDS-Defining Non-Hodgkin Lymphoma Among Carriers of HLA-B8-DR3. <i>Journal of Acquired Immune Deficiency Syndromes (1999)</i> , 2019 , 81, 266-273	3.1	1
2	African Mitochondrial DNA Haplogroup L2 Is Associated With Slower Decline of Etell Function and Lower Incidence of Diabetes Mellitus in Non-Hispanic, Black Women Living With Human Immunodeficiency Virus. <i>Clinical Infectious Diseases</i> , 2020 , 71, e218-e225	11.6	О
1	Identification of sequence variants in the CCL3 chemokine gene family in the HapMap West African reference population. <i>Human Immunology</i> , 2014 , 75, 854-8	2.3	