

Brahim Aissani

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

28

papers

871

citations

14

h-index

28

g-index

28

ext. papers

925

ext. citations

4

avg, IF

3.47

L-index

| # | Paper | IF | Citations |
|----|--|------|-----------|
| 28 | African Mitochondrial DNA Haplogroup L2 Is Associated With Slower Decline of T-cell 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 | 0 |
| 27 | 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 |
| 26 | Fine Mapping of the Body Fat QTL on Human Chromosome 1q43. <i>PLoS ONE</i> , 2016 , 11, e0153794 | 3.7 | 2 |
| 25 | Admixture mapping of genetic variants for uterine fibroids. <i>Journal of Human Genetics</i> , 2015 , 60, 533-8 | 4.3 | 10 |
| 24 | 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 | 4.8 | 17 |
| 23 | Mitochondrial DNA diversity in the African American population. <i>Mitochondrial DNA</i> , 2015 , 26, 445-51 | | 13 |
| 22 | 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 |
| 21 | 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 |
| 20 | 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 |
| 19 | A candidate gene approach for virally induced cancer with application to HIV-related Kaposi's sarcoma. <i>International Journal of Cancer</i> , 2014 , 134, 397-404 | 7.5 | 5 |
| 18 | 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 | |
| 17 | Confounding by linkage disequilibrium. <i>Journal of Human Genetics</i> , 2014 , 59, 110-5 | 4.3 | 12 |
| 16 | Mitochondrial DNA variation and virologic and immunological HIV outcomes in African Americans. <i>Aids</i> , 2014 , 28, 1871-8 | 3.5 | 1 |
| 15 | Structure of tumor necrosis factor-alpha haploblocks in European populations. <i>Immunogenetics</i> , 2013 , 65, 543-52 | 3.2 | 9 |
| 14 | Multiple hits for the association of uterine fibroids on human chromosome 1q43. <i>PLoS ONE</i> , 2013 , 8, e58399 | 3.7 | 14 |
| 13 | 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 |
| 12 | 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 |

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| 11 | 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 |
| 10 | Host genetics and HIV-1 viral load set-point in African-Americans. <i>Aids</i> , 2009 , 23, 673-7 | 3.5 | 31 |
| 9 | 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 |
| 8 | 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 |
| 7 | Compositional mapping of the human dystrophin-encoding gene. <i>Gene</i> , 1992 , 122, 329-35 | 3.8 | 28 |
| 6 | The compositional properties of human genes. <i>Journal of Molecular Evolution</i> , 1991 , 32, 493-503 | 3.1 | 61 |
| 5 | 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 |
| 4 | The distribution of genes in the human genome. <i>Gene</i> , 1991 , 100, 181-7 | 3.8 | 223 |
| 3 | CpG islands: features and distribution in the genomes of vertebrates. <i>Gene</i> , 1991 , 106, 173-83 | 3.8 | 91 |
| 2 | Compositional properties of telomeric regions from human chromosomes. <i>FEBS Letters</i> , 1991 , 295, 22-6 | 3.8 | 11 |
| 1 | CpG islands, genes and isochores in the genomes of vertebrates. <i>Gene</i> , 1991 , 106, 185-95 | 3.8 | 80 |