

List of Publications by Citations

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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.

17 papers	1,564 citations	14 h-index	17 g-index
17 ext. papers	1,729 ext. citations	12.8 avg, IF	3.44 L-index

#	Paper	IF	Citations
17	Predicting function: from genes to genomes and back. <i>Journal of Molecular Biology</i> , 1998 , 283, 707-25	6.5	370
16	Re-annotating the <i>Mycoplasma pneumoniae</i> genome sequence: adding value, function and reading frames. <i>Nucleic Acids Research</i> , 2000 , 28, 3278-88	20.1	194
15	Frameshift peptide-derived T-cell epitopes: a source of novel tumor-specific antigens. <i>International Journal of Cancer</i> , 2001 , 93, 6-11	7.5	166
14	Pathogenesis of DNA repair-deficient cancers: a statistical meta-analysis of putative Real Common Target genes. <i>Oncogene</i> , 2003 , 22, 2226-35	9.2	138
13	Secreted fringe-like signaling molecules may be glycosyltransferases. <i>Cell</i> , 1997 , 88, 9-11	56.2	112
12	HGVbase: a human sequence variation database emphasizing data quality and a broad spectrum of data sources. <i>Nucleic Acids Research</i> , 2002 , 30, 387-91	20.1	100
11	Homology-based fold predictions for <i>Mycoplasma genitalium</i> proteins. <i>Journal of Molecular Biology</i> , 1998 , 280, 323-6	6.5	100
10	Automated annotation of GPI anchor sites: case study <i>C. elegans</i> . <i>Trends in Biochemical Sciences</i> , 2000 , 25, 340-1	10.3	96
9	Systematic identification of genes with coding microsatellites mutated in DNA mismatch repair-deficient cancer cells. <i>International Journal of Cancer</i> , 2001 , 93, 12-9	7.5	83
8	HGBASE: a database of SNPs and other variations in and around human genes. <i>Nucleic Acids Research</i> , 2000 , 28, 356-60	20.1	63
7	SelTarbase, a database of human mononucleotide-microsatellite mutations and their potential impact to tumorigenesis and immunology. <i>Nucleic Acids Research</i> , 2010 , 38, D682-9	20.1	57
6	Frameshift mutations in coding repeats of protein tyrosine phosphatase genes in colorectal tumors with microsatellite instability. <i>BMC Cancer</i> , 2008 , 8, 329	4.8	24
5	Detection of coding microsatellite frameshift mutations in DNA mismatch repair-deficient mouse intestinal tumors. <i>Molecular Carcinogenesis</i> , 2015 , 54, 1376-86	5	22
4	High frequency of LMAN1 abnormalities in colorectal tumors with microsatellite instability. <i>Cancer Research</i> , 2009 , 69, 292-9	10.1	22
3	NAIL-Network Analysis Interface for Linking HMMER results. <i>Bioinformatics</i> , 2000 , 16, 656-7	7.2	7
2	Recurrent Frameshift Neoantigen Vaccine Elicits Protective Immunity With Reduced Tumor Burden and Improved Overall Survival in a Lynch Syndrome Mouse Model. <i>Gastroenterology</i> , 2021 , 161, 1288-1302.e13	13.3	6
1	(Phospho)proteomic Profiling of Microsatellite Unstable CRC Cells Reveals Alterations in Nuclear Signaling and Cholesterol Metabolism Caused by Frameshift Mutation of NMD Regulator UPF3A. <i>International Journal of Molecular Sciences</i> , 2020 , 21,	6.3	4

