Peter Beyerlein

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/11103477/publications.pdf

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10	314	7	8
papers	citations	h-index	g-index
10	10	10	907
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	mRNA-seq with agnostic splice site discovery for nervous system transcriptomics tested in chronic pain. Genome Research, 2010, 20, 847-860.	5 . 5	71
2	Sequencing of <scp>C</scp> harcot– <scp>M</scp> arie– <scp>T</scp> ooth disease genes in a toxic polyneuropathy. Annals of Neurology, 2014, 76, 727-737.	5.3	63
3	Robust gene expression and mutation analyses of RNA-sequencing of formalin-fixed diagnostic tumor samples. Scientific Reports, 2015, 5, 12335.	3.3	54
4	Targeted or whole genome sequencing of formalin fixed tissue samples: potential applications in cancer genomics. Oncotarget, 2015, 6, 25943-25961.	1.8	53
5	Association of the Charcot–Marie–Tooth disease gene ARHGEF10 with paclitaxel induced peripheral neuropathy in NCCTG N08CA (Alliance). Journal of the Neurological Sciences, 2015, 357, 35-40.	0.6	40
6	VaDiR: an integrated approach to Variant Detection in RNA. GigaScience, 2018, 7, .	6.4	16
7	Discriminative Generalized Hough transform for localization ofÂjoints in the lower extremities. Computer Science - Research and Development, 2011, 26, 97-105.	2.7	13
8	Using a priori knowledge to align sequencing reads to their exact genomic position. Nucleic Acids Research, 2012, 40, e125-e125.	14.5	4
9	Signal Processing in Genomics and Proteomics: Defining Its Role [From the Guest Editors]. IEEE Signal Processing Magazine, 2012, 29, 19-21.	5 . 6	O
10	digit—a tool fordetection andidentification ofgenomicinterchromosomaltranslocations. Nucleic Acids Research, 2017, 45, gkx010.	14.5	0