

# Peter Beyerlein

## List of Publications by Year in descending order

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

Version: 2024-02-01

10  
papers

314  
citations

1307594

7  
h-index

1588992

8  
g-index

10  
all docs

10  
docs citations

10  
times ranked

907  
citing authors

#	ARTICLE	IF	CITATIONS
1	VaDiR: an integrated approach to Variant Detection in RNA. GigaScience, 2018, 7, .	6.4	16
2	digitâ€”a tool for detection and identification of genomic interchromosomal translocations. Nucleic Acids Research, 2017, 45, gkx010.	14.5	0
3	Targeted or whole genome sequencing of formalin fixed tissue samples: potential applications in cancer genomics. Oncotarget, 2015, 6, 25943-25961.	1.8	53
4	Robust gene expression and mutation analyses of RNA-sequencing of formalin-fixed diagnostic tumor samples. Scientific Reports, 2015, 5, 12335.	3.3	54
5	Association of the Charcotâ€”Marieâ€”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	Sequencing of <scp>C</scp>harcotâ€”<scp>M</scp>arieâ€”<scp>T</scp>ooth disease genes in a toxic polyneuropathy. Annals of Neurology, 2014, 76, 727-737.	5.3	63
7	Using a priori knowledge to align sequencing reads to their exact genomic position. Nucleic Acids Research, 2012, 40, e125-e125.	14.5	4
8	Signal Processing in Genomics and Proteomics: Defining Its Role [From the Guest Editors]. IEEE Signal Processing Magazine, 2012, 29, 19-21.	5.6	0
9	Discriminative Generalized Hough transform for localization of joints in the lower extremities. Computer Science - Research and Development, 2011, 26, 97-105.	2.7	13
10	mRNA-seq with agnostic splice site discovery for nervous system transcriptomics tested in chronic pain. Genome Research, 2010, 20, 847-860.	5.5	71