

# Matias Piipari

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3967759/publications.pdf>

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6  
papers

1,158  
citations

1684188

5  
h-index

1872680

6  
g-index

6  
all docs

6  
docs citations

6  
times ranked

3959  
citing authors

#	ARTICLE	IF	CITATIONS
1	Using probabilistic estimation of expression residuals (PEER) to obtain increased power and interpretability of gene expression analyses. <i>Nature Protocols</i> , 2012, 7, 500-507.	12.0	799
2	Large-Scale Gene Regulatory Motif Discovery with NestedMICA. <i>Science, Engineering, and Biology Informatics</i> , 2011, , 3-24.	0.1	1
3	Metamotifs - a generative model for building families of nucleotide position weight matrices. <i>BMC Bioinformatics</i> , 2010, 11, 348.	2.6	6
4	iMotifs: an integrated sequence motif visualization and analysis environment. <i>Bioinformatics</i> , 2010, 26, 843-844.	4.1	11
5	The two most common histological subtypes of malignant germ cell tumour are distinguished by global microRNA profiles, associated with differential transcription factor expression. <i>Molecular Cancer</i> , 2010, 9, 290.	19.2	60
6	An ENU-induced mutation of miR-96 associated with progressive hearing loss in mice. <i>Nature Genetics</i> , 2009, 41, 614-618.	21.4	281