Jianxing Feng

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Identifying ChIP-seq enrichment using MACS. Nature Protocols, 2012, 7, 1728-1740.	12.0	1,471
2	GFOLD: a generalized fold change for ranking differentially expressed genes from RNA-seq data. Bioinformatics, 2012, 28, 2782-2788.	4.1	376
3	Using MACS to Identify Peaks from ChIPâ€Seq Data. Current Protocols in Bioinformatics, 2011, 34, Unit 2.14.	25.8	203
4	Imputation for transcription factor binding predictions based on deep learning. PLoS Computational Biology, 2017, 13, e1005403.	3.2	87
5	Inference of Isoforms from Short Sequence Reads. Journal of Computational Biology, 2011, 18, 305-321.	1.6	54