

Next-generation transcriptome assembly

Nature Reviews Genetics

12, 671-682

DOI: [10.1038/nrg3068](https://doi.org/10.1038/nrg3068)

Citation Report

#	ARTICLE	IF	CITATIONS
2	PERSPECTIVES Transcriptome sequencing: next generation approach to RNA functional analysis. <i>Biotechnologia</i> , 2011, 4, 311-319.	0.3	0
3	Next generation quantitative genetics in plants. <i>Frontiers in Plant Science</i> , 2011, 2, 77.	1.7	11
4	Alzheimer's disease models and functional genomicsâ€”How many needles are there in the haystack?. <i>Frontiers in Physiology</i> , 2012, 3, 320.	1.3	18
5	Toward 959 nematode genomes. <i>Worm</i> , 2012, 1, 42-50.	1.0	51
6	CBrowse: a SAM/BAM-based contig browser for transcriptome assembly visualization and analysis. <i>Bioinformatics</i> , 2012, 28, 2382-2384.	1.8	7
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9	RNA Sequencing: Platform Selection, Experimental Design, and Data Interpretation. <i>Nucleic Acid Therapeutics</i> , 2012, 22, 271-274.	2.0	207
10	The blood-brain barrier and pathogens. <i>Virulence</i> , 2012, 3, 157-158.	1.8	4
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15	Alternative Splicing: A Potential Source of Functional Innovation in the Eukaryotic Genome. <i>International Journal of Evolutionary Biology</i> , 2012, 2012, 1-10.	1.0	66
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18	Trinity RNA-Seq assembler performance optimization. , 2012, , .		46
19	COPE: an accurate <i>k</i>-mer-based pair-end reads connection tool to facilitate genome assembly. <i>Bioinformatics</i> , 2012, 28, 2870-2874.	1.8	145

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36	Transcriptome analysis reveals the time of the fourth round of genome duplication in common carp (<i>Cyprinus carpio</i>). <i>BMC Genomics</i> , 2012, 13, 96.	1.2	101
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