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CulebrONT: a streamlined long reads multi-assembler pipeline for prokaryotic and eukaryotic genomes

DOI: 10.24072/pcjournal.153

, , 2, .

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#	Paper	IF	Citations
4	The Complete Genome Resource of <i>Xanthomonas oryzae</i> pv. <i>oryzae</i> CIX2779 Includes the First Sequence of a Plasmid for an African Representative of This Rice Pathogen.		0
3	A strongly improved assembly of the pearl millet reference genome using Oxford Nanopore long reads and optical mapping.		0
2	A modular metagenomics analysis system for integrated multi-step data exploration.		0
1	An improved assembly of the pearl millet reference genome using Oxford Nanopore long reads and optical mapping.		0