

Computing For Comparative Microbial Genomics: Bioinformatics For Microbiologists (Computational Biology) By Trudy M. Wassenaar

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Direct repeat. Talk; Direct repeats are a type of genetic sequence that consists of two or more repeats of a specific sequence. Direct repeats are nucleotide

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The major difficulty many microbiologists face is simply that of too much information. As a result of sequencing technologies becoming so economical,

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Dr. Trudy M. Wassenaar is founder and owner of (2009) Computing for Comparative Microbial Genomics: Bioinformatics PLOS Computational Biology; PLOS

Repeated sequences (aka. repetitive elements, or repeats) are patterns of nucleic acids (DNA or RNA) that occur in multiple copies throughout the genome. The

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and innovative techniques in computational biology (bioinformatics) Computing for Comparative Microbial Genomics. Wayne, Wassenaar, Trudy M

^ Ussery, David W.; Wassenaar, Trudy; Computing for Comparative Microbial Genomics: Bioinformatics for Microbiologists. Computational Biology 8

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Definitions of repeated sequence dna, are covered in detail in "Computing for Comparative Microbial Genomics". [1] David W.; Wassenaar, Trudy;

^ a b c Ussery, David W.; Wassenaar, Trudy; Computing for Comparative Microbial Genomics: Bioinformatics for Microbiologists. Computational Biology 8

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(Redirected from Repetitive DNA) Jump to: Note: The following are covered in detail in
"Computing for Comparative Microbial Genomics". Direct repeats.