

Wjec Mark Scheme Biology By1 2014

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scientific essay from the year 2018 in the subject medicine diagnostics university of salford college of science and technology course msc biomedical science language english abstract next generation sequencing one of the applications of genomics employs bioinformatics tools and other molecular biology techniques such as the polymerase chain reaction pcr and electrophoresis to diagnose diseases once sequenced our assembled genome sequenced can be annotated to identify target genes and compared to a reference genome for similarities such as the case of blast which is used to establish evolutionary relationships between species the sequences can also be compared for variants in this report illumina based next generation sequencing was used to sequence the genomes of three staphylococcus aureus strains which are the methicillin resistant strain mrsa methicillin sensitive strain mssa and an unknown strain of s aureus u this involved an array of techniques from the extraction of genomic dna by way of the qiagen dneasy dna extraction kit to quality control of our purified dna by 1 agarose gel electrophoresis nanodrop and qubit it then moved on to preparation of dna libraries from our genomic dna extract with the qiagen qiaseq fx dna library kit from where the libraries went through an illumina miseq sequencer raw sequence data was then assembled and analysed with genome annotation tools such as atermis artemis comparison tool wgsa and abricate annotation of our s aureus species revealed antimicrobial resistance genes abricate produced 10 antimicrobial genes for mrsa 5 for mssa and 6 for u meanwhile artemis annotation gave 16 antimicrobial resistance genes for mrsa 13 for mssa and 14 for u information on antibiotic susceptibility of the three bacterial species can then be used to determine the best possible treatment for individuals with s aureus infections thus proving that next generation sequencing and genome annotation can serve as an alternative for routine disc antimicrobial susceptibility techniques to diagnose and treat infectious diseases

aggregated book

the quintessential authoritative and reader friendly rhinology reference rhinology diseases of the nose sinuses and skull base edited by distinguished clinicians peter h hwang david w kennedy richard r orlandi and peter john wormald brings together the collective wisdom experience and vision of preeminent thought leaders in a comprehensive rhinology reference with approximately 1 000 pages this tome offers in depth coverage of all core topics in the subspecialty of rhinology ranging from rhinosinusitis to endoscopic skull base reconstruction and more the updated resource reflects the significant growth in knowledge and understanding of anatomy pathophysiology medical therapies surgical techniques and technological innovations in rhinology that have occurred since the publication of the 2012 edition the second edition features a total of 69 chapters reflecting the evolving depth and breadth of the field it encompasses the pathogenesis current thinking diagnosis and management of chronic rhinosinusitis the pathology and management of less common disorders and the surgical techniques and postoperative management of skull base lesions and sinonasal tumors highlights include updated content on olfaction new material on biologic therapies enhanced coverage of office based rhinologic procedures and an

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