JAYPEE UNIVERSITY OF INFORMATION TECHNOLOGY, WAKNAGHAT TEST-3 EXAMINATION-2025

B. Tech.-V Semester (BT)

COURSE CODE (CREDITS): 18B1WBT532 (3)

MAX. MARKS: 35

COURSE NAME: COMPARATIVE AND FUNCTIONAL GENOMICS

COURSE INSTRUCTORS: DR. JATA SHANKAR

MAX. TIME: 2 Hour

Note: (a) All questions are compulsory.

(b) The candidate is allowed to make Suitable numeric assumptions wherever required

for solving problems

What are the salient characteristics of DNA Microarray? What are the basic types of DNA microarrays, and describe the major differences. Analyse the role of DNA microarray methodology to identify genome expression profiles of cancer versus non-cancerous lung cells.	CO III	5
types of DNA microarrays, and describe the major differences. Analyse the role of DNA microarray methodology to identify genome expression profiles of	i	5
role of DNA microarray methodology to identify genome expression profiles of	III	
	1 .	
If the <i>p-value</i> of differently expressed data from the Microarray is 0.01. The	CO	5
analysis enables a total of 2000 genes to be differentially expressed. What could be the estimated number of differentially expressed genes that are false	III	
Explain the working principle of Illumina sequencing technology detailing the	CO	5
major steps involved in sequencing-by-synthesis. Additionally, describe how this technology is applied in genome-wide gene expression analysis through	II	
	CO	5
humans and Saccharomyces cerevisiae. Explain the principle of gene density	CO	3
and calculate the gene density and its relationship to the biological complexity	! I	
Explain how genetic differences among individuals affect drug response. Give	CO	5
examples of diseases and drugs where genotype-guided selection or dosing is essential. Describe how pharmacogenomics supports personalised medicine in these contexts.	II	
What is a Biomarker, and describe the primary criteria for considering any	CO	5
biomolecule to be considered as a biomarker? List a few examples of cancer	III	_
P\$3 is known as the "guardian of the genome. Describe the TP53 mutations	CO	5
with example, now do these mutations contribute to cancer progression and	II	
	analysis enables a total of 2000 genes to be differentially expressed. What could be the estimated number of differentially expressed genes that are false positives or not true biological data? Explain the working principle of Illumina sequencing technology, detailing the major steps involved in sequencing-by southesis. Additionally, describe how this technology is applied in genome and gene expression analysis through RNA sequencing (RNA-seq). Evaluate and contrast the genomic architecture of eukaryotic organisms such as humans and Saccharomyces cerevisiae. Explain the principle of gene density and calculate the gene density and its relationship to the biological complexity of these species (Giventhe genomic data precisely). Explain how genetic differences among individuals affect drug response. Give examples of diseases and drugs where genotype-guided selection or dosing is essential. Describe how pharmacogenomics supports personalised medicine in these contexts. What a Biomarker, and describe the primary criteria for considering any biomolecule to be considered as a biomarker? List a few examples of cancer biomarkers? The 3 is known as the "guardian of the genome. Describe the TP53 mutations commonly observed in different types of human cancers. Discuss the concepts of loss of function, dominant-negative effects, and gain-of-function mutations.	analysis enables a total of 2000 genes to be differentially expressed. What could be the estimated number of differentially expressed genes that are false positives or not true biological data? Explain the working principle of Illumina sequencing technology, detailing the major steps involved in sequencing-by spitaesis. Additionally, describe how this technology is applied in genome wide gene expression analysis through RNA sequencing (RNA-seq). Evaluate and contrast the genomic architecture of eukaryotic organisms such as humans and Saccharomyce's celevisiae. Explain the principle of gene density and calculate the gene density and its relationship to the biological complexity of these species (Give the genomic data precisely). Explain how genetic differences among individuals affect drug response. Give examples of diseases and drugs where genotype-guided selection or dosing is essential. Describe how pharmacogenomics supports personalised medicine in these contexts. What is a Biomarker, and describe the primary criteria for considering any biomolecale to be considered as a biomarker? List a few examples of cancer biomarkers? The 3 is known as the "guardian of the genome. Describe the TP53 mutations commonly observed in different types of human cancers. Discuss the concepts of loss of function, dominant-negative effects, and gain-of-function mutations. With example, how do these mutations contribute to cancer progression and