

Paper: BI-635 Introduction to Data Analysis using R Programming

- CO1** Students will know the importance and effectiveness of R language in analysis of biological data as compared to other platforms
Activity 1 Write any five R packages and their biological relevance
Case Study Write any ten types of analyses of biological data that can be easily performed with R

Marks	Roll No	Test1	Test2	Total marks of attempted		Percentage	Score (3,2,1)	Target >= 60% (Y/N)
				questions	Marks			
			10	10	20	20		
	25101		9	9	18	20	90	3 Y
	25103		9	8	17	20	85	3 Y
	25104		8	9	17	20	85	3 Y
	25105		7	7	14	20	70	3 Y
	25106		9	10	19	20	95	3 Y
						Average	3	

%age	score
>=60	3
50-59.9	2
40-49.0	1

- CO2** Students will be able to use various R objects to handle specific type of data
T1 Explain different R objects with examples
A2 Write a note of R objects to handle sequence data

Marks	Roll No	Test 1	Assignment2	Total marks of attempted		Percentage	Score (3,2,1)	Target
				questions	Marks			
			10	10	20	20		
	25101		9	9	18	20	90	3 Y
	25103		8	8	16	20	80	3 Y
	25104		9	9	18	20	90	3 Y
	25105		7	6	13	20	65	3 Y
	25106		9	9	18	20	90	3 Y
						Average	3	

- CO3** Students can be able to read big data from local as well as remote locations
A1 Download text data with different delimiters (comma, tab, semicolon etc.) and read them using R
A2 Download data from a remote server using download.file ()function in R

Roll No	Assignment1	Assignment2	Total marks of attempted		Percentage	Score (3,2,1)	Target
			questions	Marks			

Marks		10	10		20		
	25101	8	7	15	20	75	3 Y
	25103	8	8	16	20	80	3 N
	25104	8	8	16	20	80	3 Y
	25105	6	5	11	20	55	2 N
	25106	9	8	17	20	85	3 Y
						Average	2.8

CO4 Students can be able to statistically analyze biological data

A1 Download any eukaryotic genome and count different type of nucleotides and write results to a file

A2 Download gene expression data for cyclophilin gene family in wheat and generate a heatmap using R

	Roll No	Test1	Assignment1	Total marks of attempted questions	Total Marks	Percentage	Score (3,2,1)	Target
Marks			10	10		20		
	25101	8	7	15	20	75	3 Y	
	25103	8	8	16	20	80	3 N	
	25104	8	8	16	20	80	3 Y	
	25105	6	5	11	20	55	2 N	
	25106	9	8	17	20	85	3 Y	
						Average	2.8	

CO5 Students will be able to write R scripts

T1 Write an R script to read RNAseq data, perform its Quality Control. Also draw the QC plots

A1 Using above pre-processed data generate Multidimensional scaling plots and heatmaps

	Roll No	Test1	Assignment1	Total marks of attempted questions	Total Marks	Percentage	Score (3,2,1)	Target
Marks			10	20		30		
	25101	9	18	27	30	90	3 Y	
	25103	8	15	23	30	76.66666667	3 N	
	25104	8	16	24	30	80	3 Y	
	25105	6	11	17	30	56.66666667	2 N	
	25106	9	19	28	30	93.33333333	3 Y	
						Average	2.8	