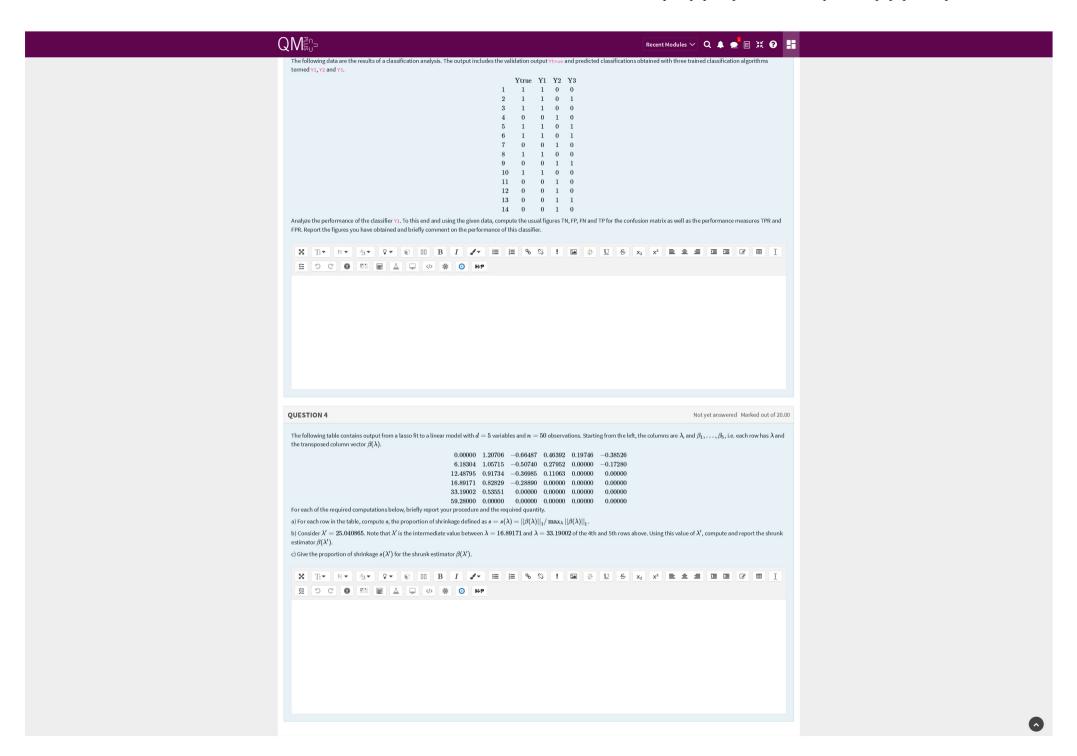
	Recent Modules 🗸 Q 🔺 🗩 💥 😧 🛔		
MTH6101 - INTRODUCTION TO MAC	HINE LEARNING - 2021/22		
Courses > Science and Engineering > MTH6101 - Intr	roduction to Machine Learning - 2021/22 > General > Semester B Final Assessment 2021/22 > Preview		
YOU CAN PREVIEW THIS QUIZ, BUT IF THIS WERI This quiz is not currently available	YOU CAN PREVIEW THIS QUIZ, BUT IF THIS WERE A REAL ATTEMPT, YOU WOULD BE BLOCKED BECAUSE: This quiz is not currently available		
QUESTION 1	Not yet answered Marked out of 20.00		
generated	variables was analysed to reduce its dimensionality. As part of Principal Component Analysis, the following variance-covariance matrix Σ was $\begin{pmatrix} 399.019 & 49.66 & -1.793 & 1.7 & 8.333 & 16.583 \\ 49.66 & 35.529 & -6.952 & 1.733 & 2.409 & 7.873 \\ -1.793 & -6.952 & 30.651 & 12.142 & -4.986 & 2.841 \\ 1.7 & 1.733 & 12.142 & 37.132 & -4.093 & -0.04 \\ 8.333 & 2.409 & -4.986 & -4.093 & 41.277 & -3.757 \\ 16.583 & 7.873 & 2.841 & -0.04 & -3.757 & 45.062 \end{pmatrix}$ ue λ_i of Σ . This eigenvalue is located in the position (4, 4) of the matrix Λ and is simultaneously the sample variance of the score PC4:		
B) Compute and write the percentage of total variability er answer.	xplained by the Principal component PC4. The number you write should be between 0 and 100 and you should include decimals in your		
C) As seen in lectures, the eigenvalue λ_4 is related to d_4 , o	one singular eigenvalue of the data matrix ${f X}.$ Compute and write the value of $d_4.$		
D) A threshold of total variability explained has been set a	t 80\%. How many principal components must you select? Write your answer.		
QUESTION 2	Not yet answered Marked out of 20.00		
	V1 V2 V3 V4 A 1.9 1.3 3.1 4.9 B 5.2 2 3.2 4.9 C 1.3 5.2 2.4 4.9 D 1.2 1 5.1 4. D 1.2 1 5.1 4. F 2.3 3.2 3.1 1.1 F 3.3 2.2 1.1 F 2.3 3.2 3.1 1.1 F 4.3 3.2 2.8 5.2 H 1.3 1.9 2.2 1.1 F 2.3 2.3 3.1.1 5.2 F 3.37 0 5.25 5.2 F 3.37 0 5.25 5.2 F 3.37 0 5.25 4.22 J 1.3 1.9 2.2 1 A B C D B F A B C D 4.22 1.9 J 4.22 2.10 5.4 4.29 3.09 D 3.375 0 5.20 4.29 3.84 J 2.32 4.24 4		
	ute and write the dissimilarity between these clusters under "average" linkage.		
C) Using the above data X, the R command XM<-kmeans (xm > KMScluster [1] 2, 1, 2, 2, 3, 3, 1, 3, 3 There is interest in determining the center of the cluster id cluster: D) Still using the above data X, the R command pam(x=X, km > PMSid.med [1] 1, 7, 6	dentified with the label 1. By computing this center manually or otherwise, identify which of the following is the correct centroid of this		
Identify correctly the medoids yielded by this cluster analy	¢		



			Recent Modules 🗸 🔍 🔺 🗩 🗄 💥 😧 🚦
Given a data set X, the library(cluster); agnes(x=X)->AG; K<-3; kmeans(x=X, ce	following ≈ commands have been run: nters=K) - >KM ojects with what you expect the ≅ output to be.		
AG\$height	Choose	÷	
AG\$order	Choose	¢	
KM\$cluster	Choose	\$	
KM\$betweenss	Choose	\$	
KM\$tot.withinss	Choose	\$	
KM\$totss	Choose	٩	
KM\$withinss	Choose	\$	
QUESTION 6			Not yet answered Marked out of 5.00
	e code is about, and what each line of code is doing. If there is outp		
QUESTION 7			Not yet answered Marked out of 5.00
O b. in some case O c. the objective	ble to use cross-validation to select a good number of clusters. s we can validate with data to determine number of clusters. is to reduce dimensionality of the data. to use cross-validation to select a good number of clusters.		
QUESTION 8			Not yet answered Marked out of 5.00
do the PCA for the san Select one: O a. reuse the eig O b. it is not possi	e performed Principal Component Analysis of a centered and unsce enset, but now centered and scaled, envalues, with the only change is to rescale them to add to the num ble to reuse eigenvalues nor eigenvectors. ted instances we can reuse eigenvalues and eigenvectors of the an	ber of components. The eigenvectors a	ance matrix to be analysed is not equal to the correlation matrix. To e the same.

