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%Part a)
T = input_data();
ClucAnalysisSingleCentroid = T(:, [4:9]);
vnames = ClucAnalysisSingleCentroid.Properties.VariableNames;
Eng = table2array(rmmmissing(T(:, [4:9])));

%Inverse-variance weights should be used for the PCA
% As the variables are in differnt units.

[wcoeff,score,latent,tsquared,explained] = pca(Eng, ...

'VariableWeights','variance');
    pc1 = score(:,1);
    pc2 = score(:,2);

    %Visuaually Showing results of PCA
    figure
    scatter(pc1, pc2, 10, 'MarkerFaceColor','blue')
    alpha(0.2)
    xlabel('1st Principal Component')
    ylabel('2nd Principal Component')
    grid()
    set(gca,'FontSize',20);

    var_by_2_first = sum(explained(1:2));
    fprintf("The first 2 principal components explain %d prct of the
variance.", var_by_2_first);
    %The first two PC account for 92% of the vareince, this close to
    %100%,this is enough.

%Part b)
figure
coef_norm = inv(diag(std(Eng)))* wcoeff;
biplot(coef_norm(:,1:2),'Scores',score(:,1:2), ...
'Varlabels',vnames);
%Interpretation:
% the direction and length of the vector indicate how each
variable
% contributes to the two principal components in the plot.

% The largest coefficients in the first principal component
% correspond to the variables `Weight` and `HorsePower`.

% The second principal component, on the vertical axis,
% has POSITIVE coefficients for the variables acceleratio, weight,
cylinders,
% and NEGATIVE coefficients for HorsePower and MPG.

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    % This indicates that the second component distinguishes among
cars
    % that have high values for the first set of variables (Weight,
Acceleration ,...)
    % and low for the second (MPG, Horsepower,...)
    % and cars that have the opposite.

%Part c)
%Classical multi-dimesnional scaling (MDS)
D = squareform(pdist(Eng));
[tmp ev] = cmdscale(D);
%Values after classical multi-dimensional scaling
% It returns 2 outputs:
% 1) the matrix tmp of the coordinates in the lower dimension
space
%    that tries to preserve the original distances
% 2) the eigenvalues (ev)  of the spectral decomposition used
inside the MDS, that
%    indicates how large the lower dimension space should be.

%Part D)
MDS = cmdscale(D, 2);
figure

%MDS with labels for manufacturer and year number
gscatter(MDS(:,1), MDS(:,2),(rmmissing(T).Mfg));
title("MDS with Manufacturer label");

figure
gscatter(MDS(:,1), MDS(:,2),(rmmissing(T).Model_Year));

title("MDS with Model Year label");

%Part E) Clustering Analysis
% We are choosing, a priori, 3 clusters in total.

[idxCluster, centroids] = kmeans(MDS,3);

figure
hold on

% Color the data points wih their respective cluster:
scatter(MDS(:,1), MDS(:,2),10, idxCluster,'Filled')
title("k-means Clustering",'FontSize',10)
alpha(0.5); grid()

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Model = string(rmmissing(T).Mfg);

%The following code will be used to label 5 points from 5
clusters
%Iterate over the cluster Variable and seperate the specific
indexis
%into the correspondign group array
%Generate 5 random numbers, to choose five random indexes for
each
%group
% Use this to annotate

indexOne = [];
indexTwo = [];
indexThree = [];
CountOne = 1;
CountTwo = 1;
CountThree = 1;
for idx = 1:numel(idxCls)

    if idxCls(idx) == 1

        indexOne(CountOne) = idx;
        CountOne = CountOne + 1;
    end

    if idxCls(idx) == 2
        indexTwo(CountTwo) = idx;
        CountTwo = CountTwo + 1;
    end

    if idxCls(idx) == 3
        indexThree(CountThree) = idx;
        CountThree = CountThree + 1;
    end

    %text(MDS(random(idx),1),MDS(random(idx),2),Model{idx} );
end
random = randi([1,80],1,5);
while(length(random) ~= length(unique(random)))

    random = randi([1,80],1,5);

end

for idx = 1:5
    text(MDS(
indexOne(random(idx)),1),MDS(indexOne(random(idx)),2),Model{idx});

text(MDS(indexTwo(random(idx)),1),MDS(indexTwo(random(idx)),2),Model{idx});

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text(MDS(indexThree(random(idx)),1),MDS(indexThree(random(idx)),2),Model{idx});
    end

    % text(MDS(random(idx),1),MDS(random(idx),2),Model{idx} );

%Part F) Clustering analysis using two

%Single Clustering
ClusterAnalysys = linkage(MDS,"single");

figure
subplot(2, 2,1);
dendrogram(ClusterAnalysys,0, 'Orientation','left')
grid()
title("Single");

ClucAnalysisSingleCentroid = cluster(ClusterAnalysys,'Maxclust',3);
subplot(2, 2,2);
scatter(MDS(:,1), MDS(:,2), 10,
ClucAnalysisSingleCentroid, 'filled')
title("Single");

%Centroid Clustering
ClusterAnalysys = linkage(MDS,"centroid");
subplot(2, 2,3);
dendrogram(ClusterAnalysys,0, 'Orientation','left')
grid()
title("Centroid");

ClucAnalysisSingleCentroid = cluster(ClusterAnalysys,'Maxclust',3);
subplot(2, 2,4);
scatter(MDS(:,1), MDS(:,2), 10,
ClucAnalysisSingleCentroid, 'filled')
title("Centroid");

    %No the clustering is not similar, and this can be attributed
to the
    %specific methods themselves. They have differnt min max
distances, thus the linkages
    %between the two are differnt, leading to clustering which is
not
    %similar.

function T = input_data()

%Function to covnert types in data table to the correct units
T = readtable("cars.csv");
T.Model_Year = double(T.Model_Year);
T.Acceleration = double(T.Acceleration);
T.Cylinders = double(T.Cylinders);

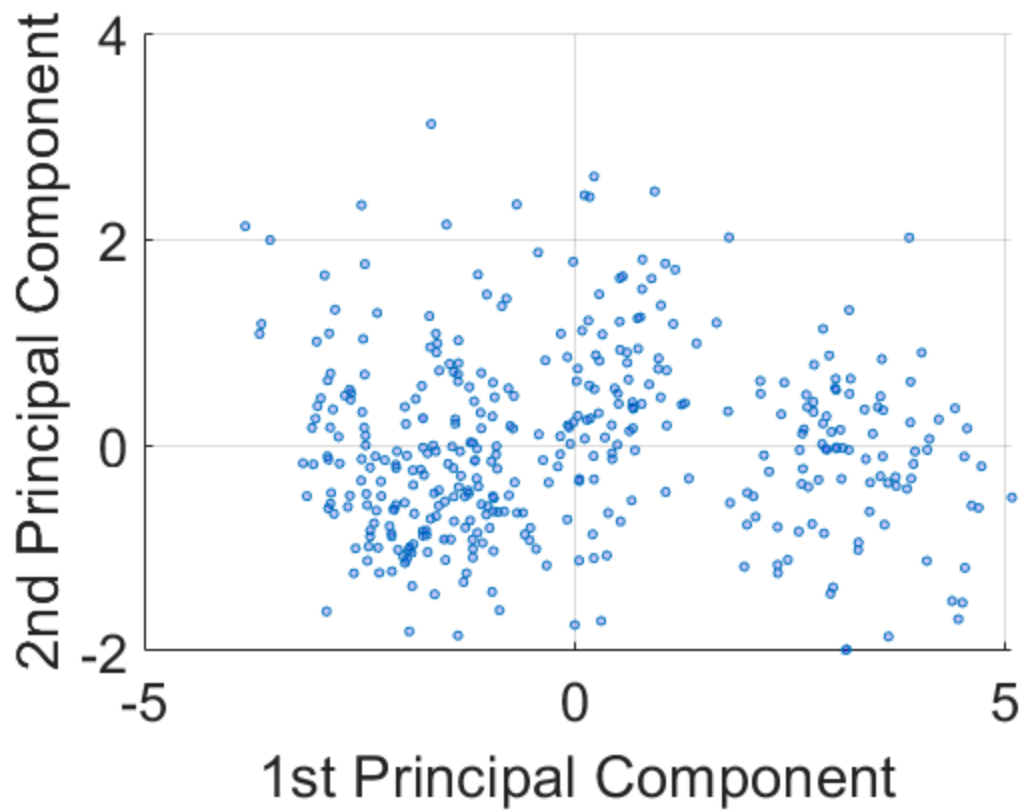
```

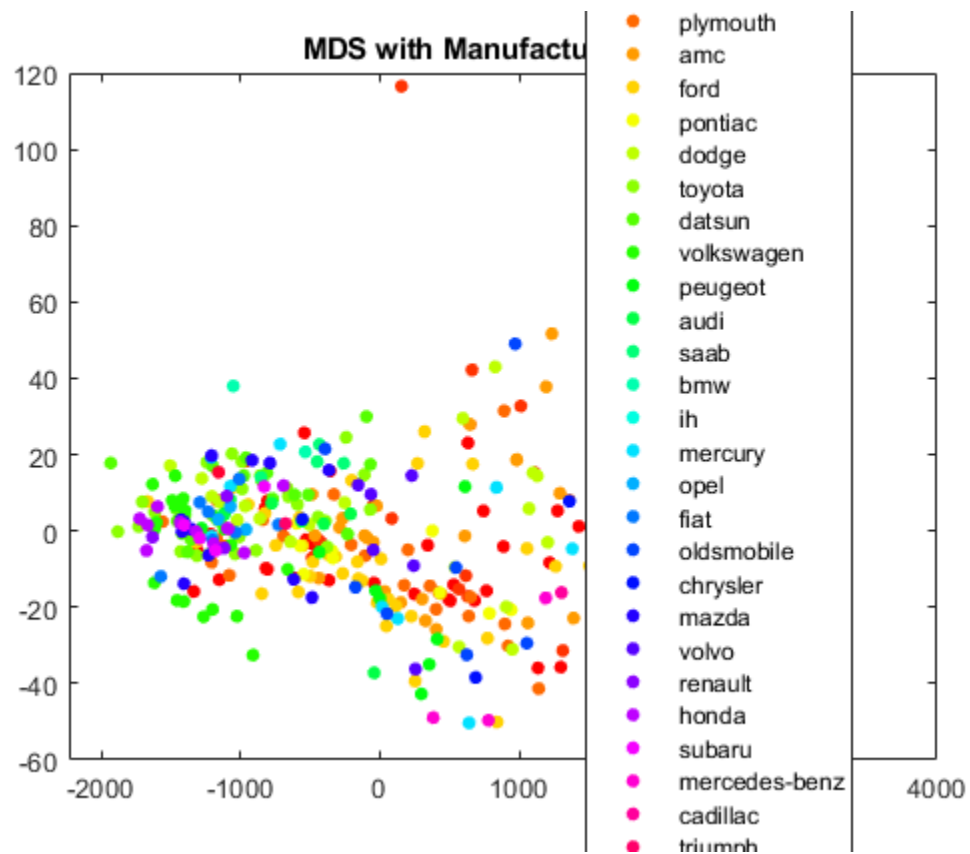
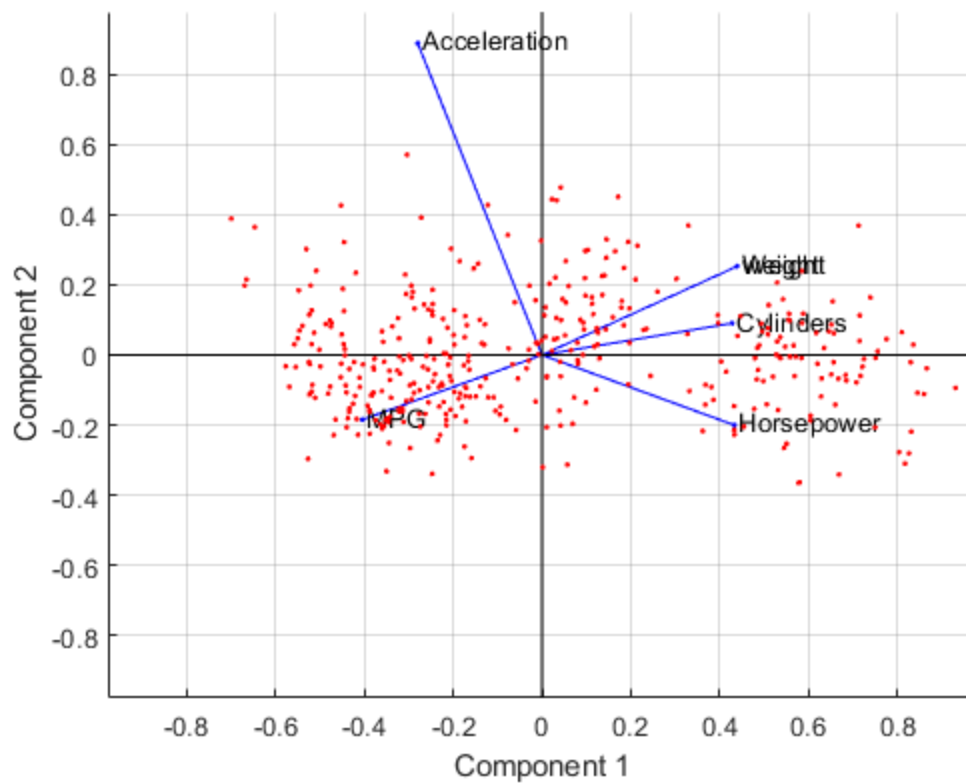
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T.Horsepower = double(T.Horsepower);  
T.MPG = double(T.MPG);  
T.weight = double(T.Weight);  
end
```

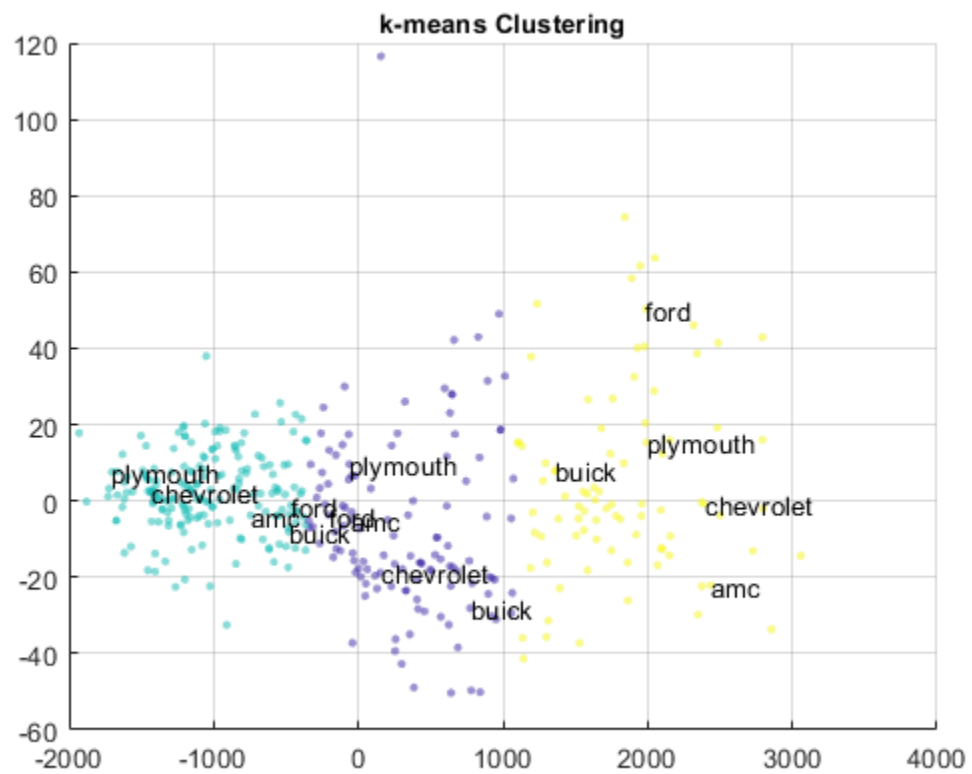
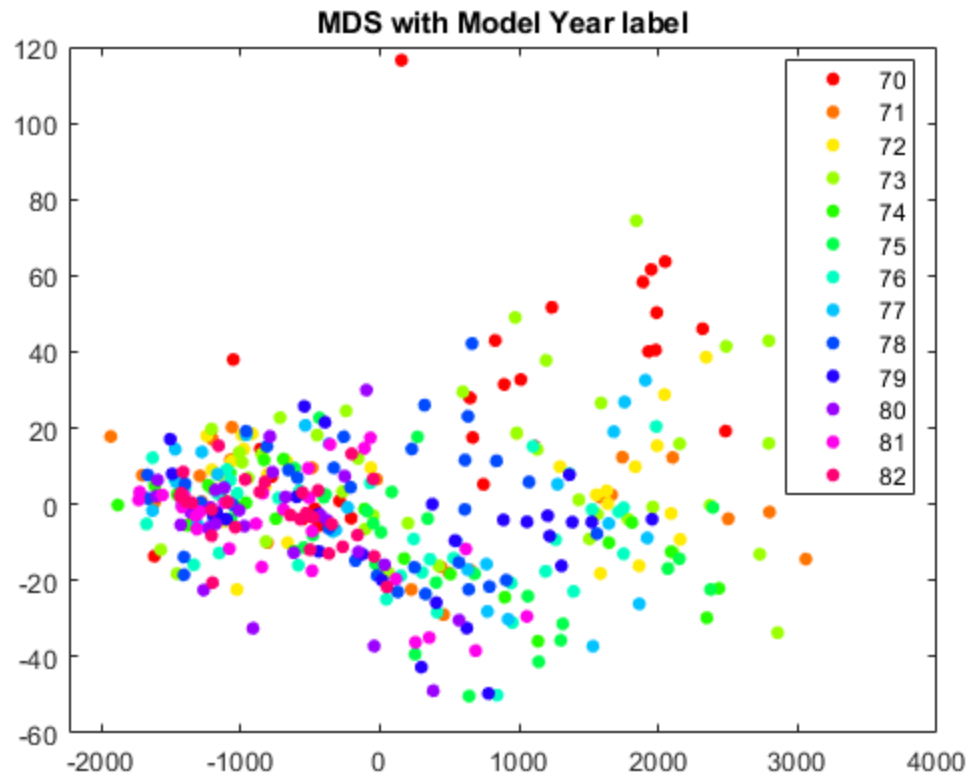
Warning: Columns of X are linearly dependent to within machine precision.

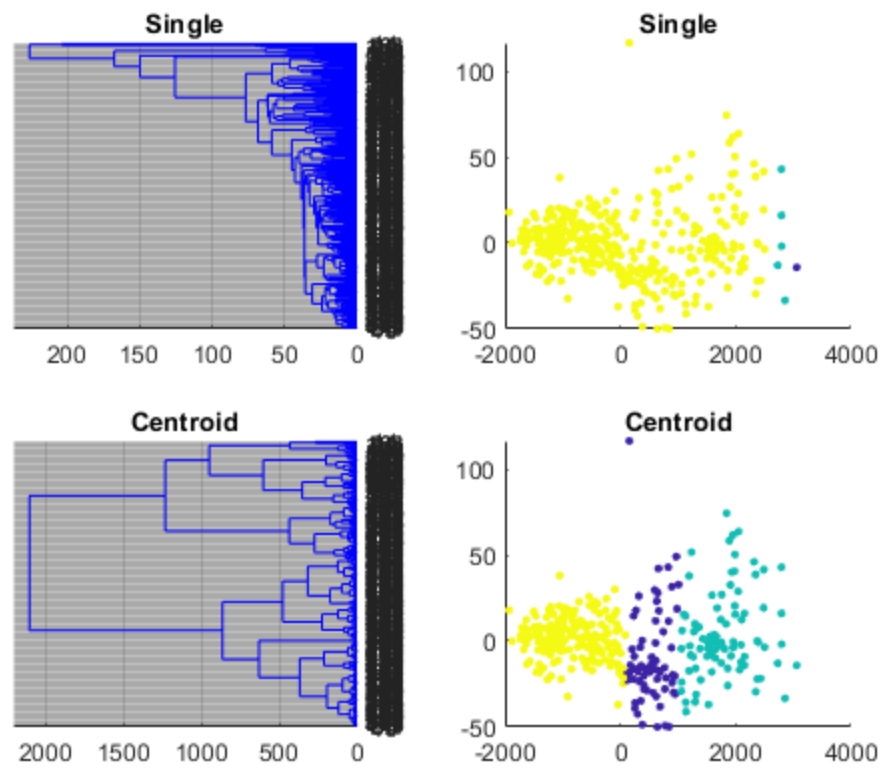
Using only the first 5 components to compute TSQUARED.

The first 2 principal components explain 9.229243e+01 prct of the variance. Warning: Non-monotonic cluster tree -- the centroid linkage is probably not appropriate.









Published with MATLAB® R2019b