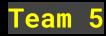
Group Assignment 3 Presentation and Demo



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Which dataset did we choose?

- Dataset Chosen: normalized weather attributes
 - This dataset contains weather attributes such as 'Air Temperature', 'Relative Humidity', 'Air Pressure', 'Track Temperature', 'Wind Speed'
 - Each feature(attribute) was normalized to ensure similar weight when clustering
- Description of features chosen
 - Air Temperature: Temperature of the air, which may influence track conditions
 - Relative Humidity: A measure of moisture in air
 - Air Pressure: Can indicate altitude or weather changes
 - Track Temperature: Surface(track) temperature, which could affect traction
 - Wind Speed: Could affect driving stability
- The dataset contains numerous data points, where each represents a unique weather update with the above attributes



K-means

Purpose of K-means

 K-means is unsupervised learning algorithm organizing data into clusters, each centered around a 'centroid'

Preparation Step

 Normalization was performed to ensure that all features are on a similar scale

How the code works for K-means

- Initialization: Randomly select k data points as the initial centroids, which represent the center of each cluster
- Assignment: Assign each data point to the nearest centroid based on Euclidean distance
- Update: Calculate the new centroids by averaging all points assigned to each cluster
- Convergence check: Repeat the previous two steps until centroids do not significantly change

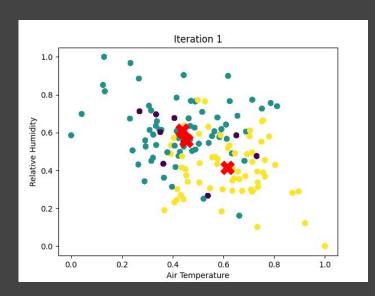


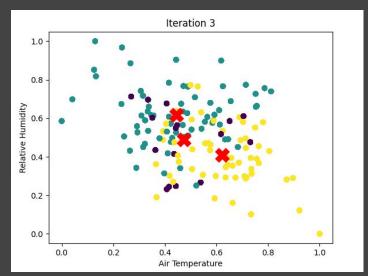
Implementation in clustering.py

- Main Functions in clustering.py
 - 1loyds: The main function conducting lloyd's algorithm for K-mans. It arranges cluster assignment and centroid updates
 - o initalizeCentroid: Initialize centroids by selecting random points from the dataset
 - assign_cluster: It calculates the Euclidean distance from each point to each centroid and assassins points to the nearest centroid
 - Calculate_centroids: The mean of points in each cluster to update the centroid positions
- Parameters used in clustering.py
 - K: Determines the number of Clusters
 - Columns: Features selected, representing weather related attributes
 - Convergence:
 - (1e-4) will stop the iterations when the movement is minimal
 - n is set to 100 by default, stopping the algorithm when it reaches
 100 iterations



Result of K-means

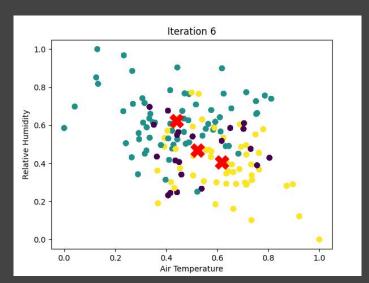


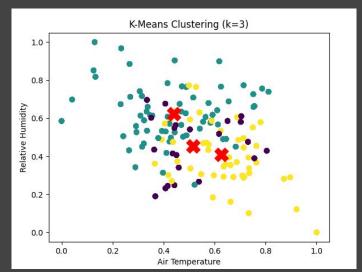


• It shows you the steps where the K-means clustering iterates by reasoning points to clusters and adjusting centroids



Result of K-means





And stops when the centroid shifts are minimal, showing the stable clusters
if max(centroid_shifts) < 1e-4:





Purpose of DBSCAN

 Unsupervised clustering method that can work well with clusters of arbitrary shape, this method model clusters as dense regions separated by sparse regions

Preparation Step

- Relevant features were normalized
- Relevant categorical features were mapped to numerical values

How the code works for DBSCAN

- Input: data points, columns, eps, min_samples
- Output: Set of density-based clusters
- Finds core points using n-dimensional Euclidean distance, n is the # of features in a data point, core point if neighbors >= min samples
- Iterates through core points, recursively adds neighboring core points to visit, loop breaks when all core points are visited



DBSCAN Implementation

```
def euclidean_distance(point1, point2):
    return math.sqrt(sum((point1[i] - point2[i]) ** 2 for i in range(len(columns))))
```

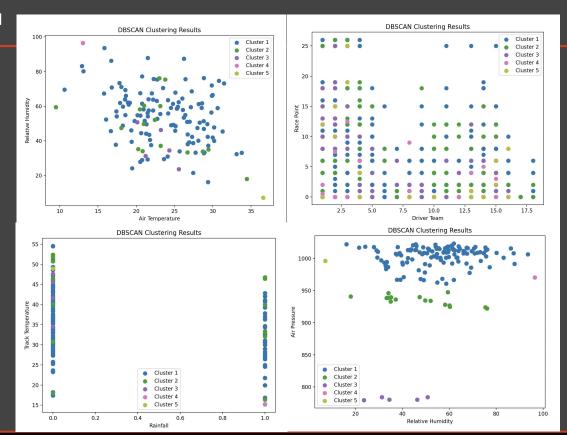
Helper functions

Recursive function loop

```
core_points = [tuple(point) for point in data if len(get_neighbors(point, data)) >= min_samples]
visited = set()
clusters = []
for core_point in core_points:
    if core_point in visited:
       continue # Skip if we've already processed this point
    cluster = []
    points_to_visit = [core_point]
       point = points_to_visit.pop()
        if point in visited:
        visited.add(point)
        cluster.append(point)
        neighbors = get_neighbors(point, data)
       if len(neighbors) >= min_samples:
           for neighbor in neighbors:
               if neighbor not in visited:
                    points_to_visit.append(neighbor)
    clusters.append(cluster) # Add the formed cluster to the list of clusters
noise = [tuple(point) for point in data if tuple(point) not in visited]
return clusters, noise
```



Result of DBSCAN





Result of DBSCAN

- Density-based clusters are formed when distances are calculated in n-dimensions
- For visualization, any two selected features may not fully represent the significance of how the clusters are formed
- Some pairs of features show better separation than others
- If trends are shown in the visuals that are not represented by the actual clusters, we can know further which features we'd want to include or exclude to show a significant separation in the visualization
 - For example, visualizing Race Points over Driver Teams
- Variance in the eps and min_samples parameters can change the silhouette score performance, occurrence of noise points, and number of clusters



Cluster evaluation - code/ approach

- In evaluating our clusters we utilized an internal method using silhouette coefficients in order to determine the accuracy of our clusters.
- The way we implemented it was through separating each part (separation point, cohesion point, and distance) into different functions
 - After the visualization and DBSCAN the math is done utilizing the formula needed to evaluate this coefficient
 - When using this we used the euclidean distance between points rather than any other distance formula.

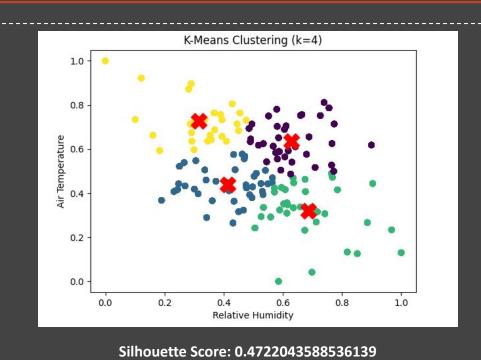


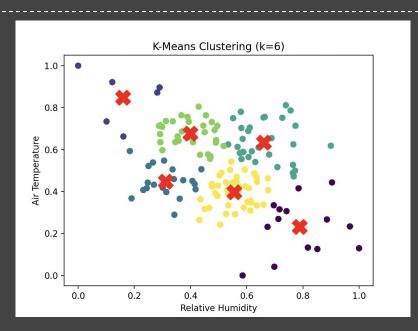
Cluster evaluation - Silhouette Score

- For this dataset, none of our group members felt confident enough with our Formula 1 knowledge to be able to produce the groundwork required for extrinsic evaluation.
- Instead, we chose to utilize the intrinsic evaluation method of the Silhouette Score.
 - The silhouette score calculates the distances of each point to the points in its cluster, as well as points in the nearest cluster.
 - For each point, the silhouette coefficient is calculated and stored.
 - At the end, the Silhouette Score is returned as the mean of the Silhouette Coefficients
- Since our K-means and DBSCAN used varying data types and methods of implementation, we decided to calculate the Silhouette scores of both these algorithms separately.
- The Silhouette Score should be between -1 and 1, where the higher the score means that points in a cluster are well matched to other points in its own cluster and poorly matched to points in other clusters.



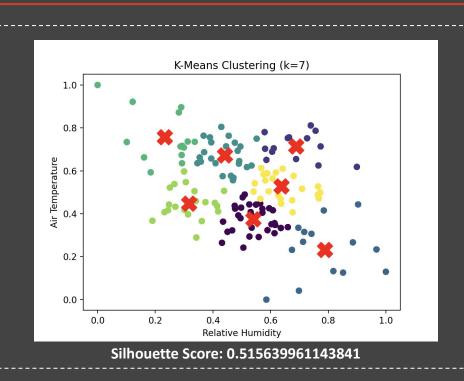
- The ideal silhouette score for k-means with our data is around
 0.7074016694527617
- This score is pretty good considering the scale of -1 to 1, so we can say that most of the points in a closer are similar to points in its own cluster, yet different from the points in different clusters
- What could be the reason for this score?
 - Looking over this there are a lot of points that do seem to skew the data away from ideal clusters
 - Since k-means uses the mean, it would make sense that outliers highly affect the clusters
 - Perhaps a better k value would increase the score
 - We'll take a look at these next





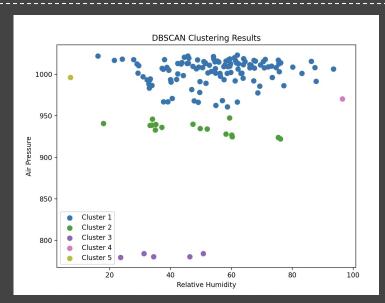
Silhouette Score: 0.6672649884420954





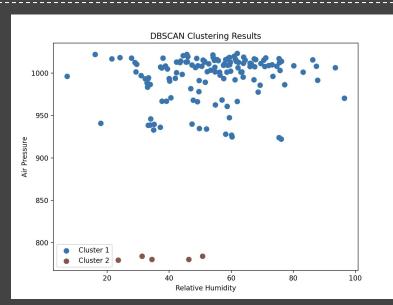


- As for DBSCAN, we calculated a Silhouette Score of around
 0.8088293062193368
- This score is pretty high considering the scaling of the Silhouette Score (recall, range of -1 to 1), and we can say that the points in the clusters are well matched
- What could be the reason for this score?
 - The radius for each core point (eps) could be optimized to include more similar points
 - The number of minimum points in each radius could be optimized in the same way
- Altering the epsilon for the DBSCAN algorithm allowed us to find the greatest Silhouette Score for our data
 - We will take a look at some examples of this



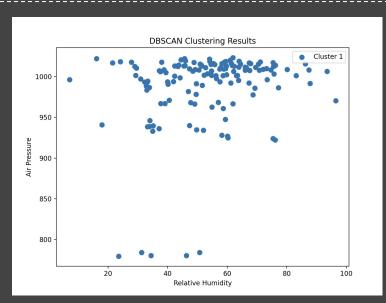
Silhouette Score: 0.36333698959419797 eps: 20





Silhouette Score: 0.8088293062193368 eps: 30





Silhouette Score: -1.0 eps: 200



Expectations vs. Reality

Expectations

 Initially expecting that normalization may not be necessary and could distort the natural structure of the data

 When looking at the scatterplot of the k-means data, we expected that the ideal k would be 4

<u>Reality</u>

- Applying normalization turned out to be important in preventing distortion, ensuring that all features contributed equally
- Our Silhouette Score with k = 4 proved this theory wrong, as we found out k = 6 produced better clusters



Issues/struggles

- The implementation of k-means proved to a lot more difficult than expected
- Issues with the actual silhouette coefficient came down to just getting a high enough score.
 - While the lower score may be errors with data collection it is something that we will have to keep in mind in order to ensure that further clustering is accurate.
- We kept encountering an issue operand type(s) for -: 'float' and 'list' with the
 data types used in dist function within testcases.py. It might be because one of
 values is a float and another one is a list
 - The issue is most likely due to mismatched data types
 - We will fix the code to access the correct level in the data structure

Thank You

