

Clustering





- Clustering uses unlabeled data and looks for similarities between groups (clusters) in order to attempt to segment the data into separate clusters.
- Keep in mind that we don't actually know the true correct label for this data!





How could we cluster this data together?

X1	X2
2	4
6	3
•••	•••
1	2

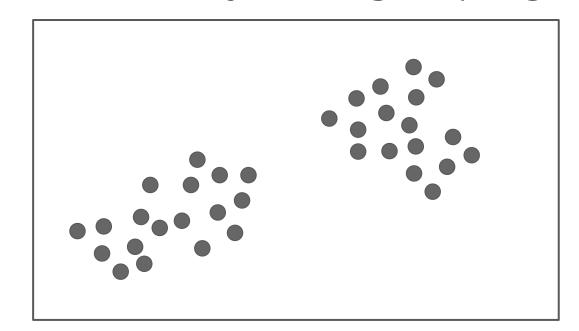




Could simply plot and discover patterns:

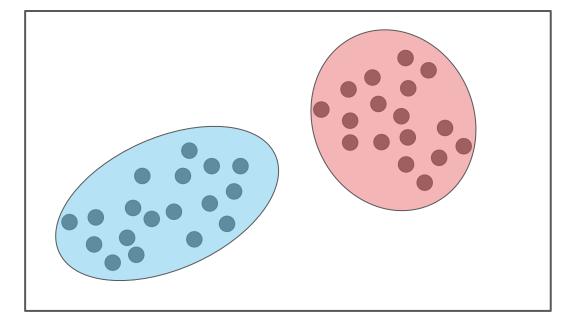


Here we intuitively see 2 groupings:



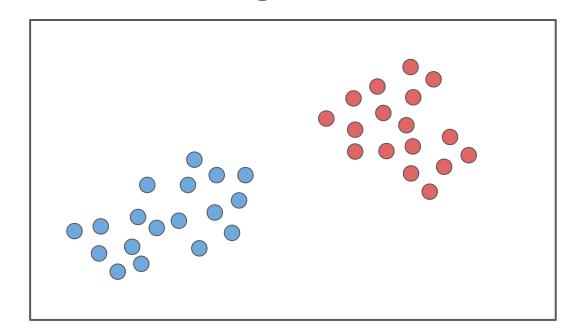


Note how distance is the intuitive metric:



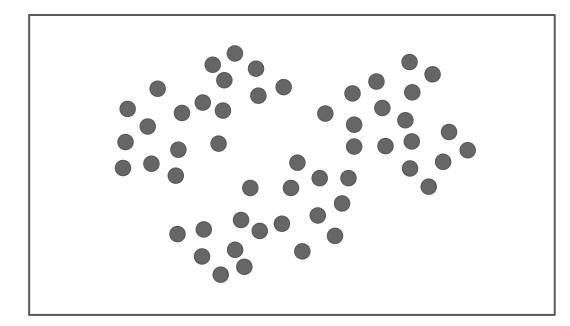


We could then assign clusters:



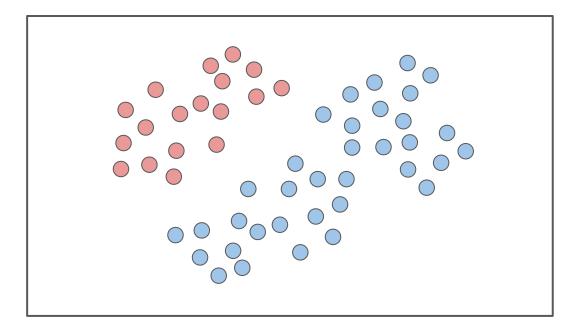


• 2 or 3 clusters could both be reasonable:



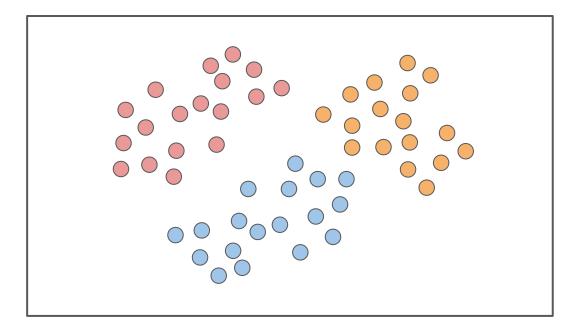


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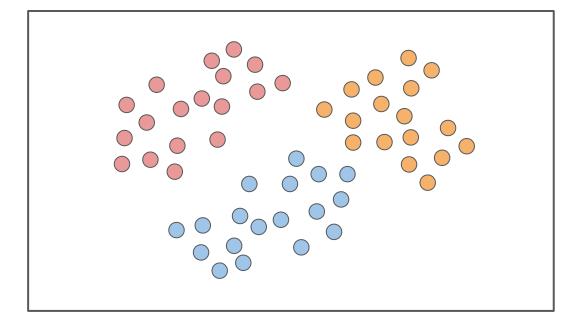


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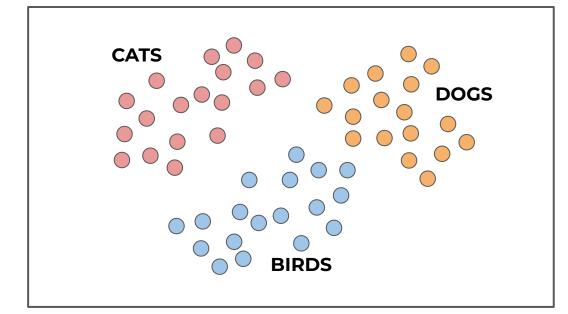


Different methods can be used to decide!





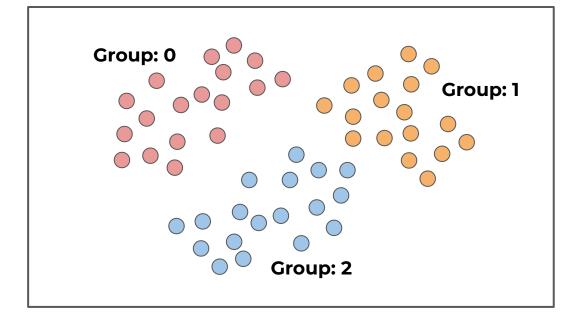
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- Unsupervised Learning Paradigm Shift:
 - If we've discovered these new cluster labels, could we use that as a y for supervised training?
 - Yes! We can use unsupervised learning to discover possible labels, then apply supervised learning on new data points.





 It's much harder to compare unsupervised algorithms against each other due to the lack of ground truth based performance metrics (e.g. can't use accuracy or RMSE).



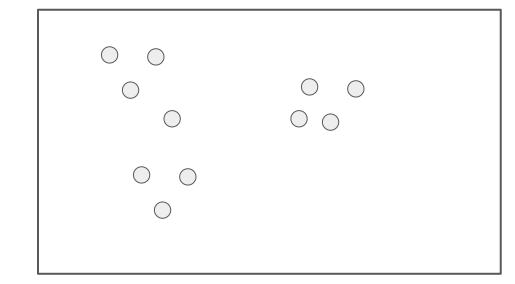


K-Means Clustering



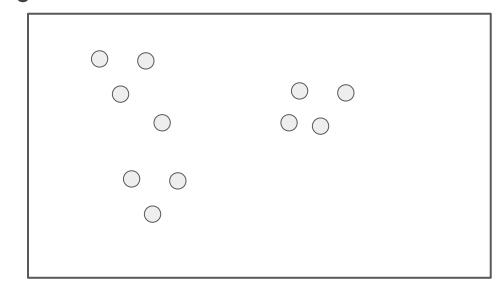


 Step 1: Choose the number of clusters to create (this is the K value).



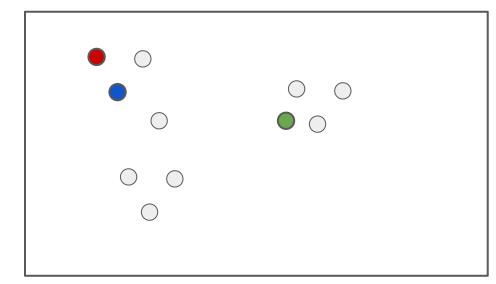


 Step 1: We'll choose K=3. Note in most situations you won't visualize the data.





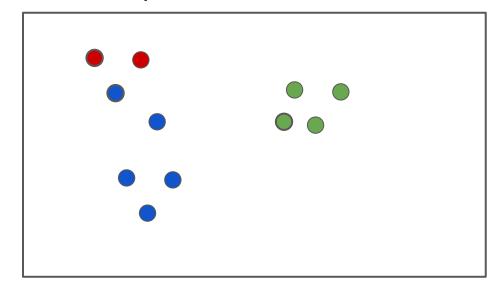
 Step 2: Randomly select K=3 distinct data points.





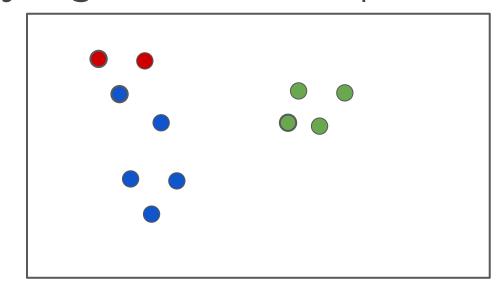


 Step 3: Assign each remaining point to the nearest "cluster" point.



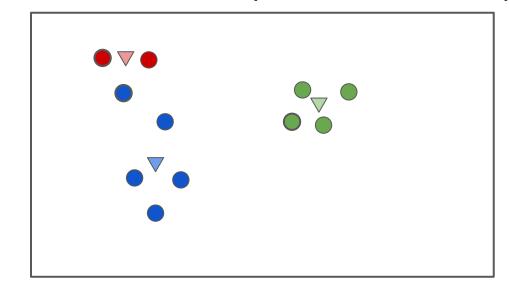


 Step 3: Note how this is using a distance metric to judge the nearest point.



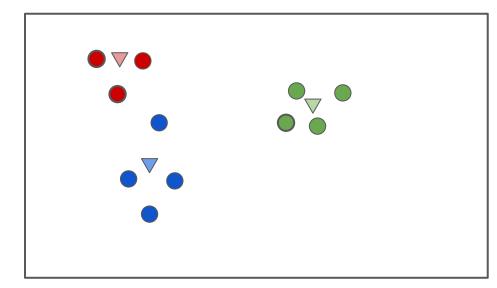


 Step 4: Calculate the center of the cluster points (mean value of point vectors).



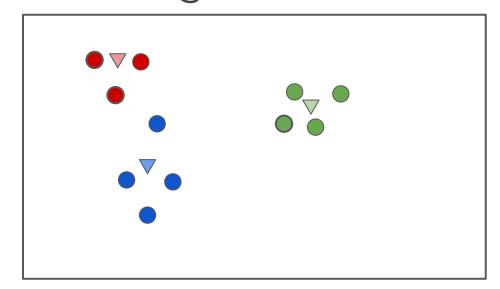


 Step 5: Now assign each point to the nearest cluster center.



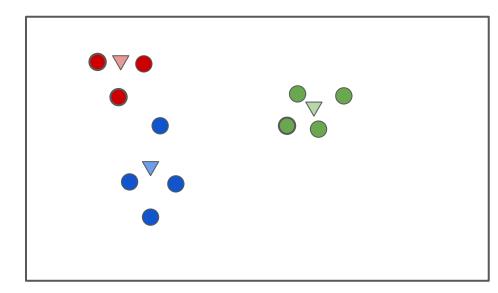


 We repeat steps 4 and 5 until there are no more cluster reassignments.





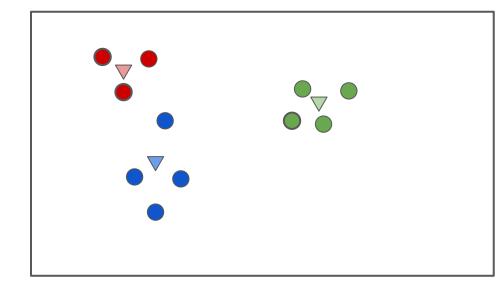
• Step 4b: Recalculate new cluster centers:







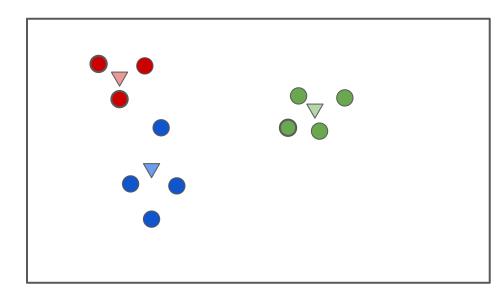
• Step 4b: Recalculate new cluster centers:





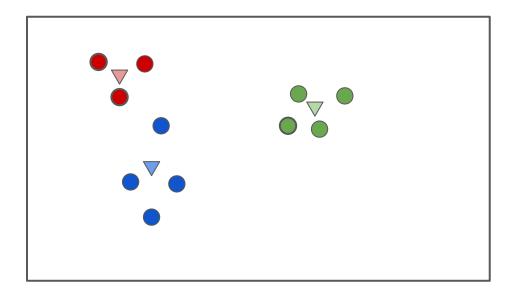


 Step 5b: Assign points to nearest cluster center.



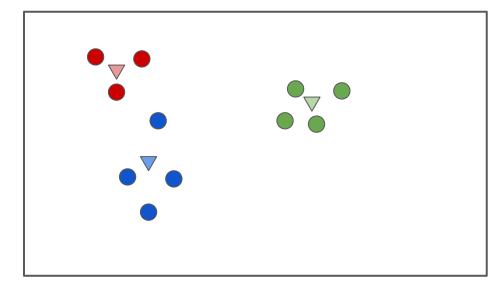


• If there are no more reassignments, we're done! The clusters have been found.



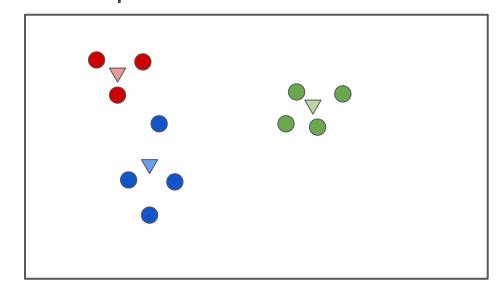


How can we measure "goodness of fit"?



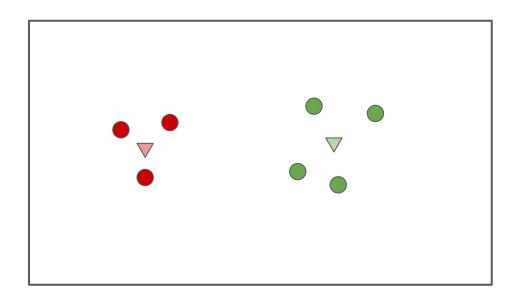


 We could measure the sum of the distances from points to cluster centers.



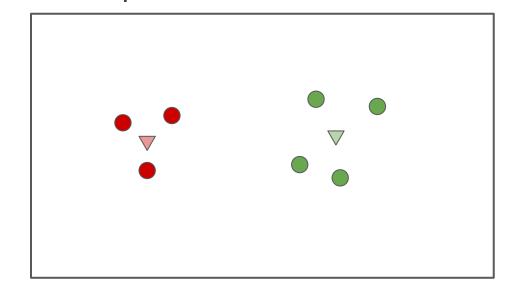


 Imagine a simple example starting with K=2.



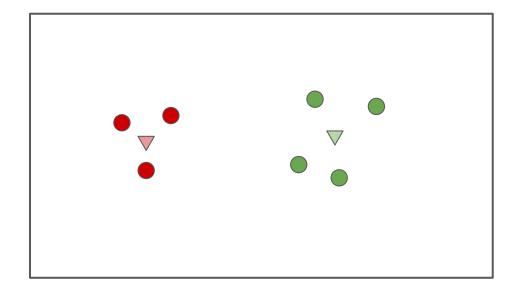


 We measure the sum of the squared distances from points to the cluster center:



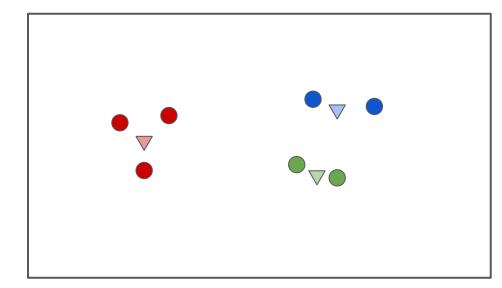


 Then we fit an entirely new KMeans model with K+1:



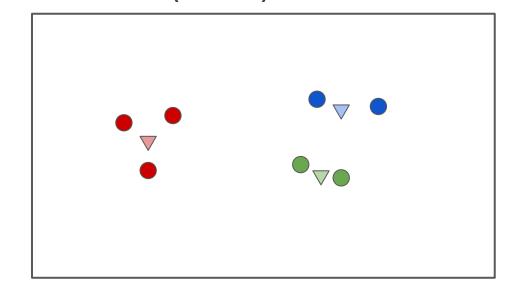


 Then we fit an entirely new KMeans model with K+1:



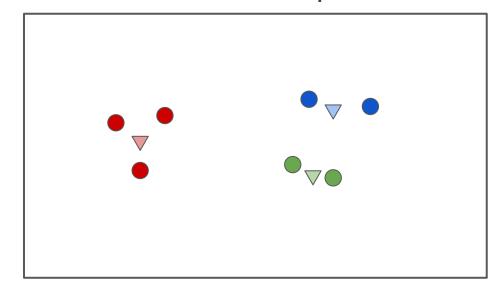


• Then measure again the sum of the squared distance (SSD) to center.



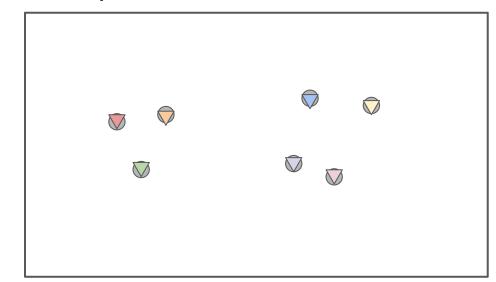


 In theory this SSD would go to zero once K is equal to the number of points.





You would have a cluster for each point!
SSD would be perfect at 0!





- We keep track of this SSD value for a range of different K values.
- We then look for a K value where rate of reduction in SSD begins to decline.
- This signifies that adding an extra cluster is not obtaining enough clarity of cluster separation to justify increasing K.



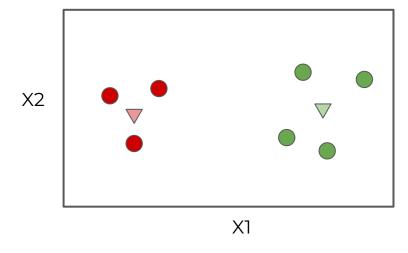


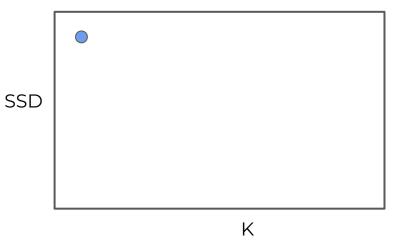
- This is known as the "elbow" method since we will track where decrease in SSD begins to flatten out compared to increasing K values.
- Let's walk through what this chart would look like...





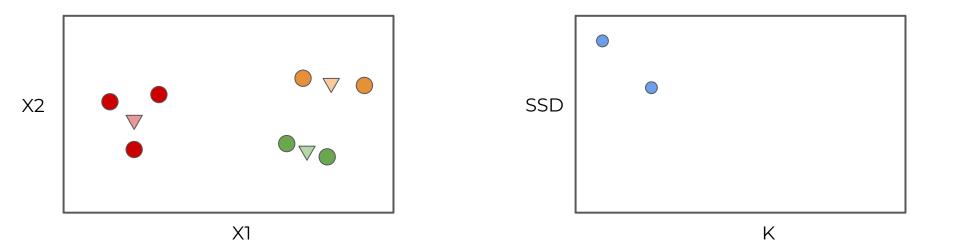
• Start with K=2:







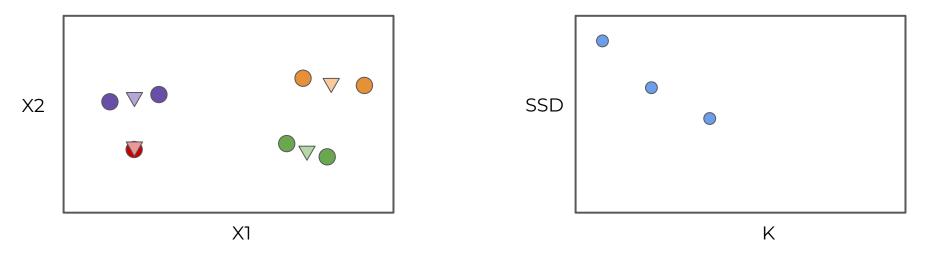
• Increase K and measure SSD:







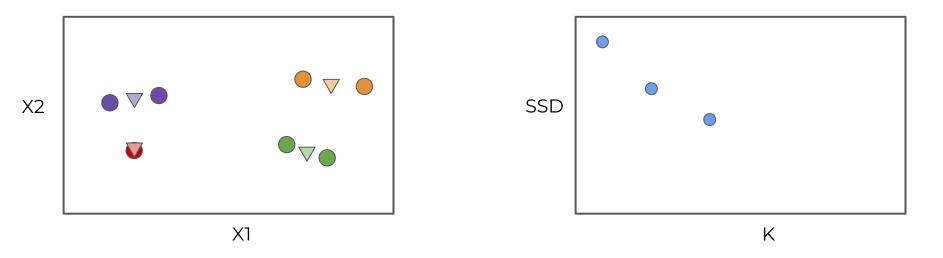
• Increase K and measure SSD:







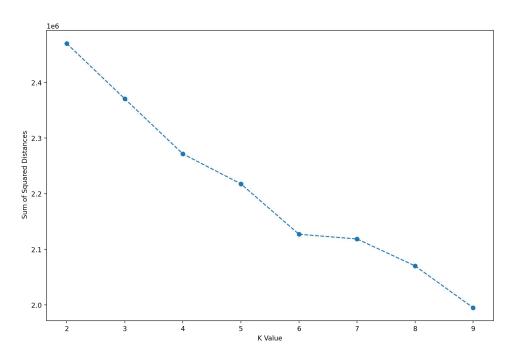
 Repeat this process for some set number of K values:







You will see a continuous decline.







- Eventually you will see "elbow" points:
- These points are strong indicators that increasing K further is no longer justified as it is not revealing more "signal".

