# Assignment #3: Unsupervised Learning and Dimensionality Reduction

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## Clustering

### Datasets

I chose to continue using my Pokemon dataset from previous assignments. This dataset contains 34 unique labels (types of Pokemon) and 25 instances of each label. The dataset is generated at runtime where each Pokemon has a type, a height, and a weight. The type for a given Pokemon is fixed while the height and weight are picked from a gaussian distribution. The means of those distributions are set in code and the standard deviation is always 10% of the mean. What is interesting about this dataset is that, given how similar some Pokemon are to others, no classifier can perfectly classify the data. This could give the clustering algorithms a tough time. I suspect that certain groups of Pokemon will get lumped into the same cluster while some Pokemon will be split among multiple clusters.

The second dataset I chose was a subset of the MNIST digit dataset. This dataset is a series of 28 pixel by 28 pixel labeled images. Each image contains a single digit. I chose to convert the images into the Weka ARFF format. The images are in greyscale which means that each entry in the dataset has 28 times 28 or 784 dimensions. I think this is a good contrast to my Pokemon dataset. The MNIST images should test the clustering algorithms ability to deal with data that has high dimensionality. The subset of the MNIST dataset I chose contains 5 digits (0-4) and 15 examples of each digit. I felt that this should be enough data to allow clustering to work while also keeping things small and easy-to-manage.

### K-Means

When using K-means clustering on the Pokémon dataset I chose a k value of 34. I figured that, since there are 34 unique labels, that made the most sense. Below is a list the predominant cluster[s] for each Pokemon.

|  |  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Bulbasaur  27 | Ivysaur  30 | Venusaur  25 | Charmander  14 | Charmeleon  32 | Charizard  22 | Squirtle  5 | Wartortle  5 | Blastoise  5 |
| Caterpie  6/8 | Metapod  17/23 | Butterfree  4/9 | Weedle  6/8 | Kakuna  7/16/23/24 | Beedrill  4/9 | Pidgey  31 | Pidgeotto  3 | Pidgeot  13/2 |
| Rattata  19 | Raticate  1/33 | Spearow  0/18 | Fearow  2/3 | Ekans  12 | Arbok  28 | Pikachu  11 | Raichu  11 | Sandshrew  21 |
| Sandslash  10/20 | Nidoran f.  15 | Nidorina  15 | Nidoqueen  28 | Nidoran m.  15 | Nidorino  15 | Nidoking  28 |  |  |

Figure 1: The predominate cluster for each Pokemon when using the K-Means method. Ideally, each Pokemon would be in its own unique cluster. For Pokemon where multiple clusters are listed, that Pokemon didn’t have a single cluster that it predominately fell into.

Looking at those clusters, I can see where clustering has issues with this kind of data. K-means did a fairly good job distinguishing between Pokemon for the most part. It stumbled exactly where I expected it to. Pokemon of the same time, with similar weights and heights tended to get grouped together into clusters that were too big. For example: because Nidoran female, Nidorina, Nidoran male, and Nidorino are all medium-sized poison Pokemon they got lumped together. There were a few surprises, however. I didn’t expect Squirtle and Blastoise to be grouped together. Although they share a type, Blastoise is 10 times heavier and 3 times taller than Squirtle. I thought that this difference would be enough to separate them. I also didn’t foresee the messy groupings for Caterpie, Metapod, Butterfree, Weedle, Kakuna, and Beedrill. These are all small bug type Pokemon, so I expected them to be grouped into a single cluster but instead they were each split among several clusters. On the whole k-means did an adequate job lining up the labels to clusters.

I chose a k value of 5 for the MNIST dataset subset because there are 5 different labels. The clusters for this data weren’t as good as I expected. In general, each label of image was placed in a unique cluster. However, cluster 0 contained too many individuals. Labels 2 and 3 were spread over several clusters. I expected that each clusters and labels would line up better than they did. I think that the issue is probably that K-Means is arriving at a stable solution too quickly. I suspect that EM will cluster this dataset much better than K-Means.

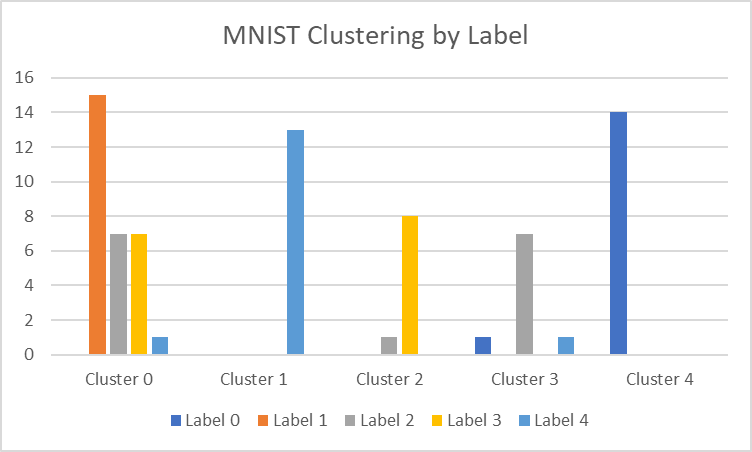


Figure 2: A bar chart indicating which clusters contained which images after running the MNIST data through a K-Means clusterer. E.g. 15 images with the label ‘0’ were put into cluster 0.

### Expectation Maximization

For both the Pokemon and MNIST datasets, I used the same k values with Expectation Maximization that with K-Means clustering. In general, EM did much better than K-Means at clustering Pokemon based on their labels. The clusters tended to be messier (i.e. a given label tended to have individuals in several clusters) but there were fewer super clusters that contain many Pokemon.

|  |  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Bulbasaur  0 | Ivysaur  15 | Venusaur  33 | Charmander  13/31 | Charmeleon  1 | Charizard  29/30/32 | Squirtle  25 | Wartortle  3/17 | Blastoise  29/32 |
| Caterpie  18 | Metapod  2 | Butterfree  22 | Weedle  16/18 | Kakuna  2 | Beedrill  22 | Pidgey  8 | Pidgeotto  11/22 | Pidgeot  5/23 |
| Rattata  7 | Raticate  20 | Spearow  14 | Fearow  23 | Ekans  12 | Arbok  28 | Pikachu  6 | Raichu  4 | Sandshrew  27 |
| Sandslash  26 | Nidoran f.  6 | Nidorina  24 | Nidoqueen  9 | Nidoran m.  13 | Nidorino  24 | Nidoking  9 |  |  |

Figure 3: The predominate cluster for each Pokemon when using the Expectation Maximization method. Ideally, each Pokemon would be in its own unique cluster. For Pokemon where multiple clusters are listed, that Pokemon didn’t have a single cluster that it predominately fell into.

As you can see the Nidoran female, Nidorina, Nidoran male, and Nidorino cluster from K-means was broken into two clusters which makes more sense given the difference in size between Nidoran male/female and Nidorina/Nidorino. Caterpie, Metapod, Butterfree, Weedle, Kakuna, and Beedrill were clustered less messily as well. To improve the quality of clustering for both EM and K-Means I would ensure that the distance metric properly handled the Pokemons’ types. Since the type attribute has no noise, any two individuals of different type should never be clumped together.

EM performed no better at clustering the MNIST data than K-Means did. This was surprising to me. I had thought that K-Means was converging too quickly because of the relatively small number of individuals in the data. I believed that EM would be able to overcome this lack of data to converge on a better set of clusters. However, if anything, EM did worse. Both zeros and twos were spread over multiple clusters. I don’t have an explanation for why this occurred.

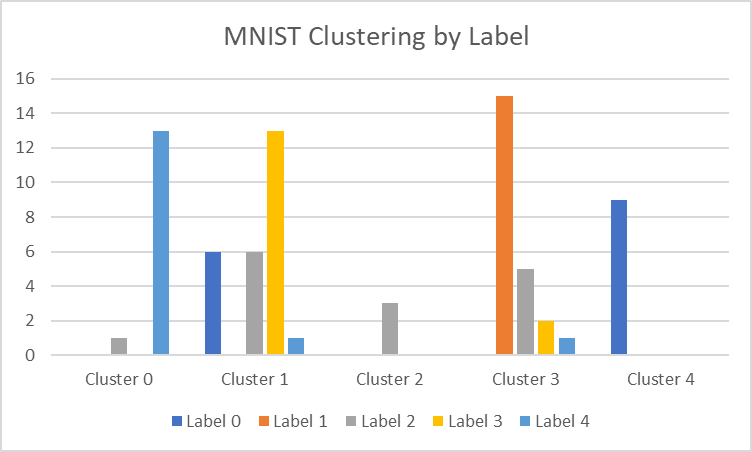


Figure 4: A bar chart indicating which clusters contained which images after running the MNIST data through an EM clusterer. E.g. 13 images with the label ‘4’ were put into cluster 0.

## Dimensionality Reduction

When I ran PCA on the Pokemon dataset, I noticed that there was very little room for dimensionality reduction. This makes sense. Since 9 out of 11 dimensions in the problem relate to the type of the Pokemon, all those dimensions are required. Combining any two of those dimensions in any way would result in a loss of information. However, weight and height are highly correlated. These two dimensions could be rolled together in one. I believe this is what I observed in the eigenvalues. There was one high eigenvalue of 2.1, one low eigenvalue of ~0, and the rest were between .2 and 1.25. The MNIST dataset is a little harder to understand. Looking at the eigenvalues, it seems like most of the information is captured in just the first 5 eigenvectors. Its also important to note that there are only 74 eigenvectors that capture any information at all (there are 75 digits in the dataset I’m using so this makes sense). I suspect that I’ll be able to get good clustering results using 5-25 eigenvectors.

## Clustering After Dimensionality Reduction

After PCA, both K-Means and EM were unable to cluster the MNIST digit dataset. There tended to be one or two large clusters that contained most of the individuals. It seems that the resulting data after being transformed by PCA is clustered in some way that doesn’t correspond to the original labels. I suspect that qualities like brightness were being selected for. This won’t work for the images because the brightness of an image is more dependent on the pen stroke width than the digit being written. The Pokemon dataset didn’t have this problem. PCA seemed to preserve most of the useful data. Doing a cursory inspection, K-Means and EM both did a little worse on the Pokemon dataset after PCA. Qualitatively, the clusters were more likely to contain multiple Pokemon and it was more likely that a single Pokemon label was split between multiple clusters.

ICA worked much better for the MNIST dataset. K-Means with ICA was a significant improvement. There were no longer problems with digits being split equally between multiple clusters. This was even more so the case for Expectation Maximization. I reran ICA -> EM several times and there tended to fewer than 6 individuals placed in the ‘wrong’ cluster (e.g. if most images with the label ‘0’ were in cluster 1 then an image with label ‘0’ placed in cluster 2 would be in the wrong cluster). Interestingly, reducing the data to 3 dimensions (from the original 784) seemed to produce the best results. That would tend to indicate that there are only a handful of vital features in the dataset. The Pokemon dataset suffered when exposed to ICA. Clusters tended to include more Pokemon. Pokemon labels often had half their individuals in one group and half in another. In general, it doesn’t seem like that dataset can be decomposed into separate components. This was true no matter how many dimensions I reduced (or even increased) the data into.