1.Introduction

I implemented PCA algorithm on my own in Java and compared the result with the result of Weka tool. I would like to share implementation of my PCA algorithm first then comparison result of it with result of Weka.

2.Implementation

2.1 Code Explanation

In this part I would like to briefly explain code. I wrote the code by looking at the notes that I took during the lectures. This is my Main, I declared all my functions as "public static" so I directly call them in my Main function.

Here is the all the variables, and functions that I declared. Data holds all the data that was read in **reader()** function. Means hold all the mean values of attributes that were calculated in **find_means()** function.

```
public static List<List<Float>> data = new ArrayList<List<Float>>();
public static List<Float> means;
public static int number_of_attributes,number_of_samples;
public static double[][] matrix; // covariance matrix
public static EigenvalueDecomposition eigenvalues;
public static Matrix eigenvector;
public static List<Integer> top_indexes=new ArrayList<>();

public static Matrix final_result;
public static int getNumber_of_attributes_reduced;
```

```
public static void find_means()
{
    means=new ArrayList<Float>();
    number_of_attributes=data.get(@).size();
    number_of_samples=data.size();
    for(int j=0; j<data.get(@).size(); j++) {
        float sum=0;
        for (int i = 0; i < data.size(); i++) {
            sum = sum + data.get(i).get(j);
        }
        means.add(sum/data.size());
    }
}</pre>
```

Reader() and **writer()** functions are easy, so I skipped that part.

In **find_means()** function I sum all attribute values of specific attribute and divide it to number of instances and store result for all attributes in **means** variable.

In the **covariance_calculator()** function I calculate covariance of any two attributes. And in **covariance_matrix()** function I create covariance matrix. In last function I find eigenvalues and eigenvectors of covariance matrix.

I used **Jama** library to find eigenvalues and eigenvectors of covariance matrix and also to do matrix operations.

```
public static float covariance_calculator(int a1, int a2)
{
    float sum=0;
    for(int i=0; i<data.size(); i++)
    {
        //(a1-mean'a1)*(a2-mean'a2)
        sum=sum+(data.get(i).get(a1)*means.get(a1) + data.get(i).get(a2)*means.get(a2));
    }
    return sum/number_of_samples;
}

public static void covariance_matrix()
{
    matrix=new double[number_of_attributes][number_of_attributes];
    for(int i=0; i<number_of_attributes; i++)
      {
            for(int j=0; j<number_of_attributes; j++)
            {
                matrix[i][j]=covariance_calculator(i,j);
            }
    }
}

public static void eigenvalue_eigenvector_finder()
{
    Matrix m=new Matrix(matrix);
        eigenvalues = m.elg();
        eigenvector=eigenvalues.getV();
}</pre>
```

Here I sort the eigenvalues that I found in previous step.

And pick the top

attribute_number_to_reduce(number of attributes in new set) .

Then I create matrix from eigenvectors that correspond to my picked eigenvalues. In this matrix each row is one eigenvector. And I multiply this matrix with my data to get new data set.

```
public static void createAmatrix_and_multiply()
{
    double A_matrix[][]=new double[top_indexes.size()][number_of_attributes];
    for(int i=0; i<top_indexes.size(); i++)
    {
        for(int j=0; j<eigenvector.getRowDimension(); j++)
        {
            A_matrix[i][j]=eigenvector.get(top_indexes.get(i),j);
        }
    }
}

    double data_matrix[][]=new double[data.size()][data.get(0).size()];

    for(int i=0; i<data.size(); i++)
    {
        for(int j=0; j<data.get(0).size(); j++)
        {
            data_matrix[i][j]=data.get(i).get(j);
        }
    }

    Matrix m1=new Matrix(A_matrix);
    Matrix m3=new Matrix(data_matrix);
    Matrix m4=m1.times(m2);
    final_result=m4.transpose();
}</pre>
```

2.2 Running the code

I also provided ReadMe file in SourceCode folder. Since I used Jama library you have to compile the code with Jama. Put your "arff" file to same folder with source code and provide name as first argument. In my code I took the location of file like this "./"+filename so putting the "arff" file to the same folder with source-code and providing name like "Data_WithoutLabels.arff" as first argument is enough. Then as second argument provide number that you want your attributes to be reduced to. It will generate reduced.arff file in the same folder, this will be your new data set with reduced number of attributes.

2.3 Testing

I reduced the number of attributes in "Data_WithoutLabels.arff" from 5 to 3 and saved it in "reduced3.arff". Then I reduced number of attributes in "reduced3.arff" from 3 to 2 and saved it in "reduced2.arff" and compared their result all of them gave same perfectly separated 6 clusters so I think my code works perfectly.

2.Comparison with Weka and result obtained.

In this section I would like to compare the result that we obtained with help of Weka and result that we obtained with my implementation.

In Result folder there exist **reduced3.arff** our new data set with 3 attribute which we obtained from our original data **Data_WithoutLabels.arff**. And also **reduced2.arff** with 2 attributes that we obtained from **reduced3.arff**. In their .xlsx files also provided below. You can see that their attributes were enough to cluster them into 6 clusters.

k	sum of squared	No	F	d				
K			X	у	Z	W	t	Instance
1	810.6925101916	1	12	6.0068	5.0143	4.002	5.0028	800
2	529.8816088737	2	5.0211	9.0041	11	9.9992	12	300
3	238.0119660525	3	6.9958	0.988	9.0006	11.007	3.008	400
4	145.8994653014	4	0.9645	2.0302	3.0318	1.0137	7.0218	100
5	75.4857954602	5	7.9877	4.01	8.0142	7.0015	9.0089	200
6	3.5446067797	6	10	7.9932	1.0127	4.9812	9.991	200
7	3.4598251315	7						
8	3.1993469413	8						
9	3.1247513846	9						
10	3.0013260538	10						
11	2.9649892881	11						
12	2.9313331396	12						
13	2.8575638188	13						
14	2.7925235051	14						
15	2.7338235933	15						

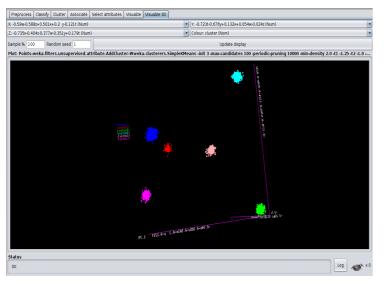
k	sum of squared	No		d				
K			X	у	Z	W	t	Instance
1	430.5284426215	1	1.443	7.079	-3.907			800
2	247.451346436	2	5.3028	18.141	-5.394			300
3	143.290481669	3	7.7874	7.6675	-1.499			200
4	37.2159727541	4	-0.778	8.9933	-10.9			400
5	19.3807744298	5		6.1416				100
6	1.7498477129	6	2.0715	11.321	-4.664			200
7	1.7126925851	7						
8	1.6571569012	8						
9	1.5689932246	9						
10	1.512211756	10						
11	1.4028013272	11						
12	1.1731091214	12						
13	1.1567508553	13						
14	1.1236086968	14						
15	1.1160063369	15						

Here the 1st result is from our original data **Data_WithoutLabels.arff** with 5 attributes, 2nd result is from our **reduced3.arff** with 3 attributes. And last one is from **reduced2.arff** with 2 attributes. I want you to pay attention to the number of instances, as you can see in all three result our total 20000 data divided to clusters as 800,300,400,200,200,100.

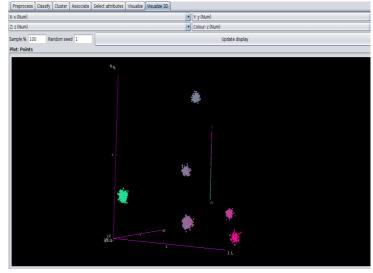
Below I provided plots for all three cases.

A	В	С	D	Е	F	G	Н	- 1	J
k	within cluster		No	Final Cluster Centroids					d
K	sum of squared		NO	X	у	Z	W	t	Instance
1	307.782522851		1	3.1407	-3.764				800
2	174.8839918157		2	9.4638	-4.983				300
3	61.2596334962		3	2.4068	-11.07				400
4	24.1883531726		4		1.5089				200
5	18.0941628601		5	4.5957	1.3813				100
6	1.4476248795		6	5.8439	-4.841				200
7	1.3381515387		7						
8	1.0724617869		8						
9	1.020239743		9						
10	0.9781228718		10						
11	0.8533624204		11						
12	0.789996291		12						
13	0.7690044855		13						
14	0.7349526715		14						
15	0.7152868286		15						

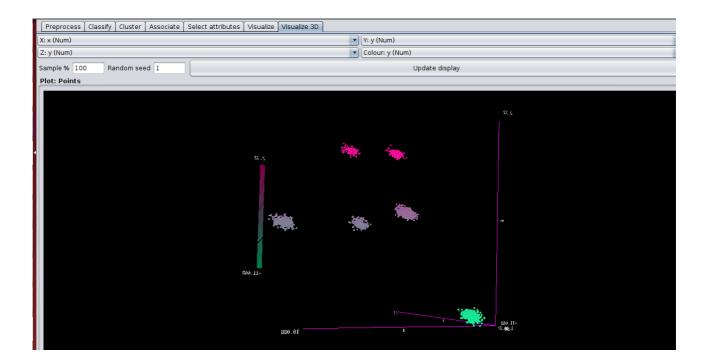
Plot of data when I reduce number of attributes to 3 with help of Weka



Plot of data when I reduce number of attributes to 3 with help of my implementation



Plot of data when number of attributes reduced to 2 with help of my implementation.



2.Conclusion

As conclusion we can clearly see from results obtained and from plots we could reduce number of attributes to 3 and to 2. This implementation was implemented with help of my lecture notes.