Machine Learning 2: Final Assignment

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1. Introduction

With the Trump administration announcing large cuts to the world's largest foreign aid donors' [6] contributions to development assistance (DA), it is clear that DA funding will decrease [12]. Given these cuts, efficiently allocating DA resources is more critical than ever to maximize impact. To increase efficiency, targeting groups of countries with similar problems beyond geographical factors can help. This report explores a key question: How can countries in need of DA be grouped based on socio-economic, demographic, health, and infrastructure indicators? My analysis of this question leverages principal component analysis (PCA) and clustering to use high dimensional data to group countries. Since development needs are characterized by many different factors, I aim to include as many variables as possible. A large problem in this kind of analysis on all countries is missing data. A single missing observation makes methods as PCA infeasible. To alleviate this I will also employ an imputation strategy, using PCA. I proceed by describing the data used in Section 2 and the method in Section 3. In Section 4, I present results and close with a discussion in Section 5.

2. Data

The data for this analysis is from the World Bank Database on World Development Indicators [1]. It contains information on 1,500 country-level development indicators. The data is extracted on all indicators for the year 2018, since it has the least missing observations. Still, around 40% observations are missing. First, I retain only one measure (e.g. when observing both % and absolute values) per indicator. Since some parts of the data have many more missing values, I drop instances where more than 10% is missing. This limits the data to around 127 variables with 192 countries observed. There are 32 demographic indicators, 51 economic, 18 geographic, 10 on health, 9 on infrastructure and 7 on regulations. The unobserved values are spread throughout the data, with mean missing per country around 1.4 (median 0), and around 1.1 (median 0) per variable. The distributions are plotted in Figure 1. Most variables have few or no observations missing, with a few outliers missing more. The distribution for missing values of countries is similar. If only a few columns or rows had many missing values, this would make imputation less suitable. A statistical summary of the variables is in Appendix Table 2. This data still has around 0.7% missing observations, spread across variables. Furthermore, for a more structured analysis of the results, I obtain data that divides countries into regions, as well as into developing and developed from the UN [11].

3. Methods

To group countries, I apply the K-Means (KM) algorithm. KM is an unsupervised machine-learning method that groups observations into clusters based on the distance between them. In high-dimensional spaces, KM's performance decreases significantly [2], making dimensionality reduction techniques such as Principal Component Analysis (PCA) useful in the case of datasets with many variables. PCA is effective in reducing the number of variables, especially when they are highly correlated. Furthermore, I leverage PCA not only for dimensionality reduction but also to impute missing values. This follows the approach by [7], who demonstrate advantages of using PCA imputed and decomposed data in KM in case of missing data. I proceed by explaining KM, followed by PCA and lastly PCA imputation.

The data is the $n \times p$ column centered matrix \mathbf{X} , with its n rows \mathbf{x}_i as row vectors of p variables observed for each individual. The KM algorithm begins by randomly assigning each observation to a cluster. The number of clusters, K, is a fixed input parameter. The centroid \mathbf{m}_k of each cluster k is then calculated as the mean of all the observations assigned to that cluster

$$\mathbf{m}_k = \frac{1}{n_k} \sum_{i=1}^{n_k} \mathbf{x}_i,$$

where \mathbf{m}_k is a vector of each of the K clusters' centroid locations. It contains the mean across the p characteristics for all assigned observations in the cluster. Using the centroids, the closest cluster for each observation is determined by

 $C_j(i) = \arg\min_k ||\mathbf{x}_i - \mathbf{m}_k||^2$

where C_j is observation i's closest centroid and $||\mathbf{x}_i - \mathbf{m}_k||^2$ the euclidean distance between observation i and a centroid k. Each observation is then assigned to the cluster whose centroid \mathbf{m}_k minimizes that distance. With this new assignment, the centroids are recalculated. This process of assignment and centroid recalculation is repeated iteratively until the cluster assignments no longer change, or until a maximum iteration criterion is reached [5]. Although euclidean distance is by far not the only distance measure, it is conventionally used when applying KM to PCA data. The results I use later also rely on this distance measure [3]. The choice of the number of clusters K can be done with data-driven methods, or with considerations from the context of the classification problem. For the data-driven methods, it is common to examine compactness of clusters in relation to the number of clusters as a diagnostic. Compactness measures how much variance is within the clusters. Since high internal similarity is desirable, this criterion serves as a diagnostic for assessing cluster quality. This is commonly evaluated using the sum of squared within-cluster distances (also known as Trace W), which should be minimized [10]. A scree plot effectively visualizes this relationship. A common approach is to identify a 'elbow' point in the graph, which serves as a good candidate for K [5].

PCA uses the singular value decomposition (SVD) to decompose the data matrix \mathbf{X} into the matrix product $\mathbf{U}\Sigma\mathbf{V}'$. \mathbf{U} is a $n\times n$ matrix of left singular vectors as columns, $\mathbf{\Sigma}$ is a $n\times p$ diagonal matrix with singular values on the diagonal, and \mathbf{V} is $p\times p$ with right singular vectors as columns. The right singular vectors are the pairwise orthogonal vectors \mathbf{v}_p that are a linear combination of the p features of the \mathbf{X} matrix. These are also called principal components (PC). The weights for this linear combination are the p elements of the vector, called loadings. The loading expresses how much of the pth feature of \mathbf{X} is used in the principal component. $\mathbf{\Sigma}$'s diagonal elements σ_p are called singular values. Their square σ_p^2 expresses the variance explained by component p. The decomposed matrices are usually order by descending variance explained. A valuable diagnostic is looking at the variance explained per component to asses how good PCA approximates \mathbf{X} . Especially looking at the cumulative sum of variance explained by the components used is informative in analyzing how well a set of dimensions does this. The left singular vectors \mathbf{u}_n are not further relevant in PCA.

Findings from [4] demonstrate that the first r PCs of the PCA decomposition are also the closest r-dimensional representation of the entire data, using a least squares error criterion. This means that the SVD restricted to the first r features is the best approximation of the p-feature original data, in the reduced

feature space. Literature [3] links the choice of the number of dimensions r to retain as input data of KM to the number of clusters K picked in KM. Specifically, they derive that r = K - 1 PCs should be used. Given we can use PCA to get matrices that closely approximate \mathbf{X} in a lower dimensional space, it is a natural choice for imputing missing data. This is done using an iterative algorithm [14]. The method consists of three steps: First, the missing values from \mathbf{X} are replaced with column means to make a first guess, producing $\hat{\mathbf{M}}_0$, a $n \times p$ matrix for imputation step 0. In the second step, PCA is applied to the data using the SVD, obtaining $\hat{\mathbf{M}}_0 = \mathbf{U} \mathbf{\Sigma} \mathbf{V}'$. Following this, we truncate the SVD to the first r < p dimensions, resulting in an close approximation of $\hat{\mathbf{M}}_0$. In the third step, the initially missing values are replaced with corresponding values from $\mathbf{U}_r \mathbf{\Sigma}_r \mathbf{V}'_r$, obtaining a new approximation matrix $\hat{\mathbf{M}}_1$. Steps 2 and 3 are then repeated until convergence, yielding the final matrix $\hat{\mathbf{M}}_c$. The initially missing values are then taken from $\hat{\mathbf{M}}_c$ as imputations in \mathbf{X} . Choosing r is crucial and can be estimated using leave one out cross-validation. This is a process searching for an optimal value of a parameter. For each candidate value of r, the process involves iteratively removing each non-missing cell, performing PCA imputation, and recording the mean squared approximation error. The total error is then summed across all simulated missing values, and the r that minimizes this error is selected as the optimal r.[8].

4. Results

The most important parameter to choose in this analysis is the number of clusters K. The goal is to create clusters that effectively distinguish multiple groups of developing countries while maintaining interpretability by limiting the number of clusters. To pick the specific value, I utilize the Scree Plot displayed in Figure 2a. It shows an elbow at 7, indicating this is a good choice for K.

I start by standardizing the data to z-scores. Following this, I impute the missing observations using PCA imputation, with an optimal r of 5 obtained via cross-validation. Then I reduce dimensionality to 6 PCs. This value is motivated by the K-1 rule from [3]. Plot Figure 2b shows that this is also a reasonable choice, as there are no strong indications to go for more or less dimensions. Using these 6 dimensions, we explain about 54% of the variance in the dataset. This is not very large, which may be a flaw of this analysis.

Using these parameters, I run the analysis. Due to the high number of variables across the 6 PCs, interpreting the 762 loadings is not feasible. However, this is also not the primary concern, as PCA is just my means for reducing data dimensionality. Nevertheless, for completeness, the loadings are included in Appendix Table 3.

As a further diagnostic, I analyze how pure the clusters are in terms of developed and developing countries. This is shown in Table 1. The results here are mixed: Three of the clusters (1, 3, 5) perfectly split off developing countries. Cluster 2 is highly mixed, while Cluster 6 consists mainly of developed countries, with three exceptions: Cuba, Armenia, and Georgia. The developed countries in this cluster are primarily from continental Europe, along with Russia and Japan. Two clusters are strongly driven by outliers: Cluster 4 isolates the United States and China, likely due to their large economies, while Cluster 7 consists solely of Djibouti. This diagnostic shows the model performance is not optimal, but I see this as acceptable

¹Note: As introduced before I am assuming a column centered **X**-matrix here. Without column centering, column means would have to be added back in step 2.

Figure 1: Histogram of Number of Missing Values

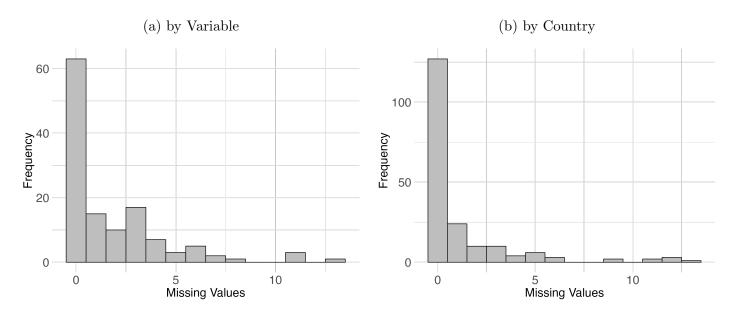


Figure 2: Scree Plots

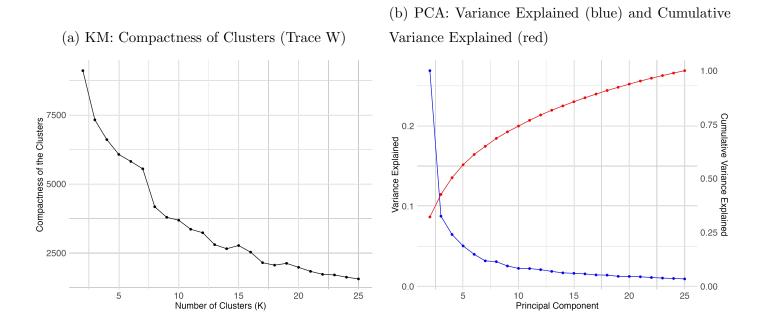


Table 1: Cluster Composition and Minority Groups

Cluster	Developed	Developing	% Developing
1	0	46	100
2	17	22	57.4
3	0	52	100
4	1	1	50
5	0	18	100
6	31	3	8.8
7	0	1	100

Note: Developed and Developing represent the number of countries with each status in each cluster.

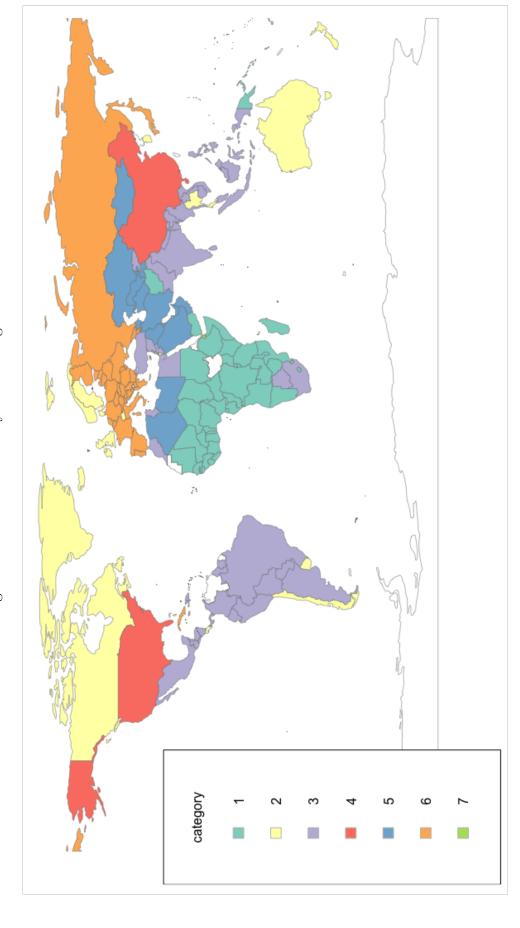


Figure 3: Countries colored by cluster assignment

for an unsupervised classification.

My main result is Figure 3, which is also represented as a table in Appendix Table 4. The map shows the assigned cluster for each country. Cluster 1 is comprised of mostly western, central and eastern African [11] countries, and additionally Yemen, Afghanistan and Papua New Guinea. The inclusion of Yemen, Afghanistan, and Papua New Guinea suggests they might benefit from similar DA strategies as these African regions. Cluster 2 is a mixed group, containing both advanced economies and smaller nations, particularly in the Caribbean. More interesting for my research question, Cluster 3 groups countries from Latin America and the Caribbean, along with regions from Southern Asia, such as India and Pakistan, Sub-Saharan Africa, Oceania, and Western Asia, including Turkey. This is the largest cluster. This cluster consists predominantly of emerging economies and aligns loosely with BRICS+ countries [13], though it notably excludes Russia, China, and some Arab states while incorporating several smaller nations. Targeting DA toward this group appears logical, as these countries share trends of rapid economic growth and more developed infrastructure. The last four clusters have less indications for DA. Cluster 4 is only comprised of the US and China. Cluster 5 groups some central Asian countries, along with Algeria and Libya and many of the large oil exporting states. Cluster 6 consists predominantly of developed economies, while Cluster 7 contains a single outlier.

5. Conclusion and Discussion

This report was able to provide interesting insights on grouping of countries based on many development indicators. The results return some groupings not conventional to DA. In summary, the analysis highlights notable groupings, such as the clustering of emerging economies, and the grouping of Western, Central, and Eastern Africa alongside Yemen, Afghanistan, and Papua New Guinea. This suggests shared development challenges and potential DA strategies. Exploring these further and analyzing synergies in providing DA based on this groups seems promising. However, the study also has some limitations:

The analysis has a very data-driven approach to selecting variables and observations. While this is sensible to reduce necessary imputations, a more theory motivated selection of variables might be better. This could significantly alter results, as it might be conceivable that variables that have more missing values across countries measure different concepts compared to more fully observed variables.

Furthermore, a diagnostic analysis on the uncertainty of the imputed data would be sensible. Using bootstrap or Bayesian methods, multiple datasets can be simulated and then analyzed. This is called multiple imputation [9]. Looking at the results from running the PCA/KM analysis could give a feeling for the uncertainty in the data due to imputation. This approach is outlined in [14]. An interesting expansion would be the use of the data in multi-dimensional scaling. The first few PCs could be used to construct a dissimilarity matrix, possibly further including geographic distance. Using this, the same research question could be examined from a different angle. Exploring these avenues could lead to further new groupings of countries based on development indicators. Testing the implementation of these groups would provide interesting insights on increasing efficiency in DA to aid as many people as possible.

References

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Appendix

Table 2: Descriptive Statistics Summary

Variable	Mean	SD	Min	Max
Access to electricity (% of population)	84.9193	24.8232	9.5000	100.0000
Adolescent fertility rate (births per 1,000 women ages	46.7664	40.1103	1.1790	165.5440
15-19) Age dependency ratio (% of working-age population)	E0 0000	17.0814	17 7015	104.0599
Agricultural land (% of land area)	$58.9080 \\ 37.7554$	22.2938	$17.7845 \\ 0.5233$	$104.9582 \\ 81.3421$
Agriculture, forestry, and fishing, value added (% of	10.2754	9.6842	0.0223	39.0099
GDP)				
Aquaculture production (metric tons)	5.4881e + 05	4.9507e + 06	-1.9930e+06	6.6135e+07
Arable land (% of land area)	14.5282	13.7238	0.0863	59.7100
Bird species, threatened	23.1146	25.7732	0	175.0000
Birth rate, crude (per 1,000 people)	20.0509	9.9958	6.4000	46.1270
Capture fisheries production (metric tons)	4.9436e+05	1.4965e+06	0	1.4831e+07
Carbon dioxide (CO2) emissions (total) excluding	760.2778	4.4853e + 03	-72.2038	4.7109e + 04
LULUCF (% change from 1990) Compulsory education, duration (years)	9.8472	2.3988	0	16.0000
Control of Corruption: Estimate	-0.0400	0.9721	-1.7858	2.1718
Death rate, crude (per 1,000 people)	7.6314	2.7000	0.9820	15.4000
DEC alternative conversion factor (LCU per US\$)	994.5485	4.7830e + 03	0.3020	5.4219e + 04
Energy intensity level of primary energy (MJ/ $$2017$	4.6702	2.6157	0.4800	18.5800
PPP GDP)	400 8004	10.000		4.42.2222
Export unit value index $(2015 = 100)$ Export value index $(2015 = 100)$	$109.5964 \\ 135.4763$	$10.0029 \\ 183.9540$	71.6000 33.6169	143.3000 $2.6366e+03$
Export value index $(2015 = 100)$ Export volume index $(2015 = 100)$	124.1947	176.1307	27.5866	2.5035e+03 2.5035e+03
Fertility rate, total (births per woman)	2.6582	1.3264	0.9770	7.0230
Fish species, threatened	39.0938	38.8610	0	251.0000
Forest area (% of land area)	32.3271	24.2310	0	94.8290
Forest rents (% of GDP)	1.3615	2.8288	0	20.7466
GDP (constant 2015 US\$) GDP deflator (base year varies by country)	4.2627e+11 217.1417	1.7996e + 12 602.2174	8.1762e+07 30.7319	1.9652e+13 6.1818e+03
,				
GDP growth (annual %) GDP per capita (constant 2015 US\$)	3.2491 $1.4719e+04$	2.5399 $2.0187e+04$	-6.2365 265.6724	8.7762 $1.0684e+05$
GDP per capita growth (annual %)	2.0207	2.7027	-9.3704	8.3850
Government Effectiveness: Estimate	-0.0374	0.9518	-2.2737	2.2318
Import unit value index $(2015 = 100)$	107.8656	4.8716	84.4000	123.4000
Import value index $(2015 = 100)$	118.4693	31.7053	33.0400	413.8667
Import volume index $(2015 = 100)$	110.0140	29.3419	33.1143	389.6674
Incidence of tuberculosis (per 100,000 people) Industry (including construction), value added (% of	109.7145	158.5430	0	1.1800e+03
,,,	25.5988	11.5246	4.0907	65.8764
GDP) Land area (sq. km)	6.6732e + 05	1.8456e + 06	20.0000	1.6377e + 07
Life expectancy at birth, female (years)	75.1126	7.7976	52.7700	87.6100
Lower secondary school starting age (years)	11.8194	0.8621	10.0000	14.0000
Mammal species, threatened	17.1615	22.4977	0	191.0000
Merchandise exports (current US\$) Merchandise exports by the reporting economy (current	9.9635e+10 9.8471e+10	2.7465e+11 2.7486e+11	-3.9097e+10	2.4867e+12 2.5013e+12
US\$)	3.04116+10	2.14008+11	1.3153e + 07	2.50158+12
Merchandise exports to economies in the Arab World	7.5047	13.4004	0.0020	75.3253
(% of total merchandise exports)	1.0041	10.4004	0.0020	10.0200
Merchandise exports to high-income economies (% of	57.5884	24.3564	2.9664	95.7271
total merchandise exports) Merchandise exports to low- and middle-income	13.6993	18.9415	0.0370	93.0144
economies in East Asia & Pacific (% of total	19.0339	10.9410	0.0910	JJ.U144
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merchandise exports)				

Table 2: Descriptive Statistics Summary (continued)

Variable	Mean	SD	Min	Max
Merchandise exports to low- and middle-income	5.8258	11.0948	0	64.8180
economies in Europe & Central Asia (% of total				
merchandise exports) Merchandise exports to low- and middle-income	4.5164	9.3840	1.0000e-04	64.3281
economies in Latin America & the Caribbean (% of				
total merchandise exports)				
Merchandise exports to low- and middle-income	2.8115	6.8815	0	64.4347
economies in Middle East & North Africa (% of total				
merchandise exports) Merchandise exports to low- and middle-income	5.9395	13.1499	0	94.5910
economies in South Asia (% of total merchandise				
exports) Merchandise exports to low- and middle-income economies in Sub-Saharan Africa (% of total	7.4657	15.0695	0.0010	92.0587
merchandise exports) Merchandise exports to low- and middle-income	22.0377	19.2953	0.1403	85.3454
economies outside region (% of total merchandise				
exports) Merchandise imports (current US\$)	1.0126e+11	2.8964e+11	-4.3613e+10	2.6142e + 12
Merchandise imports by the reporting economy (current	1.0081e+11	2.8646e+11	9.9027e+07	2.5427e + 12
US\$) Merchandise imports from economies in the Arab World	5.7792	8.7617	4.0000e-04	58.0877
(% of total merchandise imports) Merchandise imports from high-income economies (% of	55.9546	20.7732	6.8969	95.8614
total merchandise imports) Merchandise imports from low- and middle-income	17.9500	12.9462	1.4195	89.3399
economies in East Asia & Pacific (% of total	11.0000	12.0102	1.1100	00.5500
merchandise imports) Merchandise imports from low- and middle-income	7.2663	11.6880	0.0011	66.6218
economies in Europe & Central Asia (% of total				
merchandise imports)				
Merchandise imports from low- and middle-income	4.7202	8.0762	0.0018	41.6579
economies in Latin America & the Caribbean (% of				
total merchandise imports) Merchandise imports from low- and middle-income	1.6315	2.4062	0	17.1454
economies in Middle East & North Africa (% of total				
merchandise imports) Merchandise imports from low- and middle-income	4.2144	8.9654	0.0307	90.9473
economies in South Asia (% of total merchandise				
imports) Merchandise imports from low- and middle-income	6.3135	13.9392	0.0040	84.4018
economies in Sub-Saharan Africa (% of total				
merchandise imports) Merchandise imports from low- and middle-income	25.1477	15.1640	0.6812	70.0032
economies outside region (% of total merchandise				
imports)				
Merchandise trade (% of GDP) Methane (CH4) emissions (total) excluding LULUCF	$63.6231 \\ 74.7990$	$40.4956 \\ 383.7868$	13.8501 -78.2349	330.3758 $5.2244e+03$
(% change from 1990) Mineral rents (% of GDP) Mortality rate, adult, female (per 1,000 female adults) Natural gas rents (% of GDP)	0.6221 122.0313 0.8883	1.6224 81.8364 3.8607	$0 \\ 23.0550 \\ 0$	9.6111 419.4490 48.1395

Table 2: Descriptive Statistics Summary (continued)

Variable	Mean	SD	Min	Max
Net barter terms of trade index (2015 = 100) Net migration Nitrous oxide (N2O) emissions (total) excluding	103.0247 6.3676e+03 51.7933	10.7003 2.2226e+05 79.1771	53.5375 -1.3096e+06 -47.8954	136.9241 1.7650e+06 554.5455
LULUCF (% change from 1990) Official exchange rate (LCU per US\$, period average) Oil rents (% of GDP)	824.5127 2.9529	3.6830e+03 7.7308	0.3020 0	4.0864e+04 45.7659
People practicing open defecation (% of population) People using at least basic drinking water services (% of	$6.7947 \\ 88.3125$	$12.2354 \\ 15.5556$	$0 \\ 35.8471$	$68.4855 \\ 104.8187$
population) People using at least basic sanitation services (% of	76.8181	27.2296	8.3499	100.0000
population) Permanent cropland (% of land area) Plant species (higher), threatened	4.4590 85.4167	6.9561 194.6361	0.0035	39.5062 $1.8590e+03$
PM2.5 air pollution, mean annual exposure	25.8375	17.2759	5.4970	93.3070
(micrograms per cubic meter) Political Stability and Absence of Violence/Terrorism:	-0.0616	0.9756	-2.9962	1.9368
Estimate Population ages 0-14 (% of total population) Population ages 00-04, female (% of female population) Population ages 05-09, female (% of female population)	27.6346 9.5581 9.2065	10.5863 4.2531 3.5135	11.5797 3.4664 3.6236	49.0475 19.0975 16.4440
Population ages 10-14, female (% of female population) Population ages 15-19, female (% of female population) Population ages 15-64 (% of total population) Population ages 20-24, female (% of female population) Population ages 25-29, female (% of female population)	8.4804 7.9530 63.6225 7.7903 7.7716	2.8819 2.2871 6.5435 1.6501 1.3137	3.3214 3.6032 48.7904 4.1731 4.5792	13.6503 12.0170 84.9008 10.5950 12.8862
Population ages 30-34, female (% of female population) Population ages 35-39, female (% of female population) Population ages 40-44, female (% of female population) Population ages 45-49, female (% of female population) Population ages 50-54, female (% of female population)	7.3990 6.7264 6.0169 5.5457 5.1408	1.2918 1.1492 1.2128 1.4612 1.7368	5.0520 3.7818 3.0049 2.8685 2.1391	14.5421 11.8534 10.7131 8.9058 9.1678
Population ages 55-59, female (% of female population) Population ages 60-64, female (% of female population) Population ages 65 and above (% of total population) Population ages 65-69, female (% of female population) Population ages 70-74, female (% of female population)	4.6635 3.9525 8.7429 3.2023 2.3764	1.9214 1.9295 6.2440 1.8705 1.5919	1.7520 1.3073 1.1443 0.7104 0.4298	8.4412 7.8092 28.3452 7.6865 6.9113
Population ages 75-79, female ($\%$ of female population) Population ages 80 and above, female ($\%$ of female	$1.8063 \\ 2.4104$	$1.4000 \\ 2.3489$	$0.0803 \\ 0.1861$	$6.0596 \\ 10.8385$
population) Population density (people per sq. km of land area) Population growth (annual %) Population, female	347.8778 1.2026 1.9645e+07	1.6329e+03 1.2786 7.1436e+07	0.1365 -2.8988 5.6030e+03	2.0027e+04 3.9802 6.8566e+08
PPP conversion factor, GDP (LCU per international \$) Preprimary education, duration (years) Price level ratio of PPP conversion factor (GDP) to	343.1077 4.0395 0.5538	1.5197e+03 1.6418 0.2423	0.1885 1.0000 0.1741	1.6923e+04 7.0000 1.3167
market exchange rate Primary education, duration (years) Primary school starting age (years)	5.7745 6.0571	0.8417 0.5336	4.0000 5.0000	8.0000 7.0000
Regulatory Quality: Estimate Renewable energy consumption (% of total final energy	-0.0154 30.4714	0.9548 27.9458	-2.2152	2.2213 96.4000
consumption) Rule of Law: Estimate Rural population Rural population growth (annual %)	-0.0405 1.7847e+07 0.1247	0.9555 $7.8948e+07$ 1.5996	-2.2987 0 -5.0191	2.0344 9.0686e+08 3.3612
Scientific and technical journal articles Secondary education, duration (years) Secure Internet servers Services, value added (% of GDP) Sex ratio at birth (male births per female births)	1.3603e+04 6.4168 2.4108e+05 56.3585 1.0516	5.3523e+04 0.8926 1.6037e+06 11.6358 0.0200	-6.5627e+03 4.0000 -2.3493e+05 33.1147 1.0100	5.3230e+05 9.0000 2.1517e+07 94.2951 1.1480

Table 2: Descriptive Statistics Summary (continued)

Variable	Mean	SD	Min	Max
Surface area (sq. km) Survival to age 65, female (% of cohort)	7.2145e+05 80.1483	2.1082e+06 11.9293	20.0000 44.2315	1.7098e+07 95.9002
Terrestrial and marine protected areas (% of total	11.9930	11.9295 12.3285	4.0000e-04	82.9424
territorial area) Terrestrial protected areas (% of total land area) Total fisheries production (metric tons)	16.8313 1.0353e+06	11.7510 6.2171e+06	0.0492 -2.3287e+06	53.6230 8.0966e+07
Total greenhouse gas emissions excluding LULUCF (%	125.5981	412.9404	-68.4883	5.3127e + 03
change from 1990) Total greenhouse gas emissions excluding LULUCF per	6.8950	8.7096	0.0574	72.2634
capita (t CO2e/capita) Total natural resources rents (% of GDP) Tuberculosis case detection rate (%, all forms) Urban population	5.9843 76.1777 2.1723e+07	9.8486 15.3305 7.4136e+07	0 23.0000 1.1477e+04	62.7690 110.0000 8.2976e+08
Urban population growth (annual %) Voice and Accountability: Estimate	1.8498 -0.0155	$\begin{array}{c} 1.6356 \\ 0.9638 \end{array}$	-2.7136 -2.1529	5.8587 1.7093

Table 3: PCA Loadings

Variable Cumulative Variance Explained	PC1 (26.89%)	PC2 (35.64%)	PC3 (42.1%)	PC4 (47.13%)	PC5 (51.12%)	PC6 (54.29%)	PC7 (57.36%)
Access to electricity (% of population) Adolescent fertility rate (births per 1,000	-0.137 0.142	0.009 -0.002	-0.122 0.076	0.015 -0.043	-0.067 0.05	-0.016 -0.061	$0.082 \\ 0.058$
women ages 15-19) Age dependency ratio (% of working-age	0.133	-0.005	0.18	0.024	0.086	0.041	0.029
population) Agricultural land (% of land area) Agriculture, forestry, and fishing, value	0.034 0.131	-0.036 -0.017	0.101 0.076	$0.152 \\ 0.065$	-0.099 -0.041	0.079 -0.039	0.046 0.045
added (% of GDP)							
Aquaculture production (metric tons) Arable land (% of land area) Bird species, threatened Birth rate, crude (per 1,000 people)	-0.016 -0.015 0.004 0.164	-0.226 -0.015 -0.187 -0.008	-0.002 0.129 -0.033 0.053	0.017 0.151 -0.017 -0.01	-0.069 -0.087 -0.047 0.038	-0.047 0.011 -0.158 0.022	-0.006 0.034 0.049 -0.004
Capture fisheries production (metric tons)	-0.026	-0.253	-0.004	-0.005	-0.049	-0.081	-0.004
Carbon dioxide (CO2) emissions (total)	0.009	0.017	-0.022	-0.051	0.03	-0.069	-0.026
excluding LULUCF (% change from 1990) Compulsory education, duration (years) Control of Corruption: Estimate Death rate, crude (per 1,000 people) DEC alternative conversion factor (LCU	-0.063 -0.122 -0.035 0.022	0.008 0.033 0.018 -0.041	-0.01 0.063 0.233 -0.084	-0.016 -0.174 0.159 0.066	-0.049 0.027 0.104 0.037	0.017 0.041 -0.067 -0.129	0.14 -0.105 -0.003 -0.053
per US\$)							
Energy intensity level of primary energy	0.059	-0.025	-0.015	0.012	0.123	-0.06	-0.064
(MJ/\$2017 PPP GDP) Export unit value index (2015 = 100) Export value index (2015 = 100) Export volume index (2015 = 100) Fertility rate, total (births per woman)	-0.003 0.013 0.014 0.156	0.021 0.006 0.006 -0.009	-0.071 0.009 0.017 0.075	0.09 0.079 0.072 0.002	0.257 -0.129 -0.149 0.073	-0.011 0.133 0.134 0.048	-0.076 -0.3 -0.293 -0.01
Fish species, threatened Forest area (% of land area) Forest rents (% of GDP) GDP (constant 2015 US\$) GDP deflator (base year varies by	0.006 -0.014 0.1 -0.045 0.034	-0.199 0.009 -0.006 -0.246 -0.008	0.009 0.051 0.104 0.053 -0.023	-0.046 -0.101 -0.008 -0.051 0.047	-0.022 0.075 0.069 0.042 0.038	-0.044 -0.25 -0.047 0.091 0.165	0.055 0.006 -0.058 0.003 0.168
country)							
GDP growth (annual %) GDP per capita (constant 2015 US\$) GDP per capita growth (annual %) Government Effectiveness: Estimate	0.013 -0.109 -0.046 -0.142	-0.034 0.012 -0.018 0.001	0.037 0.007 0.041 0.025	0.111 -0.14 0.135 -0.125	-0.171 0.102 -0.199 0.036	-0.111 0.118 -0.13 0.009	-0.151 -0.109 -0.108 -0.102

Table 3: PCA Loadings (continued)

Import unit value index $(2015 = 100)$	-0.06	0.057	0.105	0.076	0.083	-0.011	-0.074
Import value index $(2015 = 100)$	-0.028	0.01	0.06	0.11	-0.198	0.006	-0.298
Import volume index $(2015 = 100)$ Incidence of tuberculosis (per $100,000$	-0.019 0.088	0.003 -0.021	$0.043 \\ 0.032$	0.099 -0.036	-0.209 0.002	0.007 -0.152	-0.291 -0.094
people) Industry (including construction), value	0.027	-0.063	-0.183	0.058	0.182	-0.066	-0.06
added (% of GDP) Land area (sq. km)	-0.017	-0.197	-0.008	0	0.076	0.014	0.036
Life expectancy at birth, female (years) Lower secondary school starting age	-0.156 0.052	0.009 -0.001	-0.07 0.081	-0.018 -0.192	-0.007 0.034	0.035 -0.006	0.031 -0.033
(years) Mammal species, threatened Merchandise exports (current US\$) Merchandise exports by the reporting	0.03 -0.067 -0.067	-0.178 -0.237 -0.237	-0.029 0.046 0.051	0.002 -0.033 -0.032	-0.058 0.053 0.052	-0.18 0.075 0.074	0.005 -0.048 -0.045
economy (current US\$)							
Merchandise exports to economies in the	0.035	0.002	-0.023	0.054	-0.071	0.261	0.035
Arab World (% of total merchandise							
exports) Merchandise exports to high-income	-0.075	0.014	0.074	-0.041	-0.05	-0.017	0.049
economies (% of total merchandise							
exports) Merchandise exports to low- and	0.029	-0.036	-0.104	-0.051	0.126	-0.157	-0.103
middle-income economies in East Asia &							
Pacific (% of total merchandise exports) Merchandise exports to low- and	-0.029	0.028	-0.006	0.261	0.048	-0.009	0.097
middle-income economies in Europe &							
Central Asia (% of total merchandise							
exports) Merchandise exports to low- and	-0.016	-0.008	-0.018	-0.106	-0.115	-0.024	0.217
middle-income economies in Latin							
America & the Caribbean (% of total							
merchandise exports)							
Merchandise exports to low- and	0.005	0.008	-0.031	0.076	-0.058	0.232	0.131
middle-income economies in Middle East							
& North Africa (% of total merchandise							
exports) Merchandise exports to low- and	0.05	-0.011	-0.015	0.004	0.031	0.045	-0.07
middle-income economies in South Asia							
(% of total merchandise exports) Merchandise exports to low- and	0.079	0.007	0.101	-0.043	-0.057	0.069	-0.139
middle-income economies in Sub-Saharan							
Africa (% of total merchandise exports) Merchandise exports to low- and	0.017	-0.035	-0.067	-0.007	0.136	0.171	-0.081
middle-income economies outside region							
(% of total merchandise exports) Merchandise imports (current US\$)	-0.065	-0.238	0.062	-0.047	0.053	0.092	-0.034
Merchandise imports by the reporting	-0.066	-0.238	0.063	-0.046	0.051	0.091	-0.034
economy (current US\$) Merchandise imports from economies in	0.059	-0.044	-0.064	0.054	-0.052	0.254	-0.052
the Arab World (% of total merchandise							
imports)							

Table 3: PCA Loadings (continued)

variable	PC1	PC2	PC3	PC4	PC5	PC6	PC7
Merchandise imports from high-income	-0.1	0.046	0.042	-0.09	-0.004	0.058	0.003
economies (% of total merchandise							
imports) Merchandise imports from low- and	0.041	-0.071	-0.092	-0.025	-0.004	-0.187	-0.084
middle-income economies in East Asia $\&$							
Pacific (% of total merchandise imports) Merchandise imports from low- and	-0.014	0.026	-0.042	0.291	0.089	0.005	0.062
middle-income economies in Europe $\&$							
Central Asia (% of total merchandise							
imports)							
Merchandise imports from low- and	-0.008	-0.017	-0.045	-0.102	-0.085	-0.02	0.25
middle-income economies in Latin							
America & the Caribbean (% of total							
merchandise imports) Merchandise imports from low- and	0.035	-0.041	-0.024	0.141	-0.007	0.265	0.034
middle-income economies in Middle East							
& North Africa (% of total merchandise							
imports) Merchandise imports from low- and	0.048	-0.003	-0.007	0.01	-0.051	0.019	-0.111
middle-income economies in South Asia							
(% of total merchandise imports) Merchandise imports from low- and	0.082	0.004	0.097	-0.064	0.027	0.019	-0.09
middle-income economies in Sub-Saharan							
Africa (% of total merchandise imports) Merchandise imports from low- and	0.039	-0.068	-0.041	0.047	0.076	0.202	-0.038
middle-income economies outside region							
(% of total merchandise imports)							
Merchandise trade (% of GDP) Methane (CH4) emissions (total)	-0.046 0.036	0.048 -0.001	-0.012 -0.046	0.056 -0.047	$0.013 \\ 0.126$	-0.01 -0.015	-0.251 -0.014
excluding LULUCF (% change from 1990) Mineral rents (% of GDP) Mortality rate, adult, female (per 1,000	$0.04 \\ 0.145$	-0.017 -0.005	-0.014 0.1	0.061 -0.03	0.066 0.016	-0.134 -0.032	-0.074 -0.047
female adults) Natural gas rents (% of GDP)	0.014	0.002	-0.102	-0.01	0.092	-0.065	-0.017
Net barter terms of trade index (2015 $=$	0.03	0	-0.145	0.053	0.247	-0.006	-0.015
100) Net migration Nitrous oxide (N2O) emissions (total)	-0.036 0.079	-0.055 -0.016	0.058 -0.147	-0.082 -0.056	0.149 -0.02	$0.077 \\ 0.051$	0.024 -0.076
excluding LULUCF (% change from 1990) Official exchange rate (LCU per US\$,	0.018	-0.045	-0.094	0.062	0.023	-0.166	-0.064
period average) Oil rents (% of GDP)	0.028	-0.013	-0.21	0.026	0.215	0.067	-0.026
People practicing open defecation (% of	0.11	-0.017	0.085	-0.016	0.03	-0.022	-0.068
population) People using at least basic drinking water	-0.143	0.02	-0.102	-0.015	-0.054	0.003	0.053
services (% of population) People using at least basic sanitation	-0.146	0.019	-0.11	0.023	-0.031	0.035	0.06
services (% of population) Permanent cropland (% of land area) Plant species (higher), threatened	0.017 0.013	0.024 -0.132	0.025 -0.006	-0.067 -0.042	-0.158 -0.047	-0.075 -0.106	0.086 0.077
PM2.5 air pollution, mean annual	0.09	-0.044	-0.087	0.101	-0.024	0.133	-0.07
exposure (micrograms per cubic meter)	-			-			•
1 (

Table 3: PCA Loadings (continued)

variable	PC1	PC2	PC3	PC4	PC5	PC6	PC7
Political Stability and Absence of	-0.107	0.068	0.025	-0.173	0.009	-0.097	-0.116
Violence/Terrorism: Estimate							
Population ages 0-14 (%)	0.166	-0.009	0.043	-0.019	0	0.017	0.026
Population ages 00-04, female (% f)	0.165	-0.009	0.012	-0.018	0.036	0.042	-0.003
Population ages 05-09, female (% f)	0.165	-0.007	0.007	-0.032	0.007	0.042	0.011
Population ages 10-14, female (% f)	0.163	-0.007	0.018	-0.054	-0.041	0.027	0.028
Population ages 15-19, female (% f)	0.155	-0.007	0.007	-0.077	-0.098	0.001	0.045
Population ages 15-64 (%) Population ages 20-24, female (% f)	-0.127 0.132	0.004 -0.002	-0.193	-0.031 -0.105	-0.078 -0.136	-0.026 -0.023	-0.046 0.024
Population ages 25-29, female (% f)	0.132 0.062	-0.002 -0.005	-0.084 -0.225	-0.103 -0.099	-0.130 -0.127	-0.023	-0.054
Population ages 30-34, female (% f)	-0.008	-0.005	-0.279	-0.063	-0.079	-0.007	-0.089
Population ages 35-39, female (% f)	-0.081	0.001	-0.239	-0.04	-0.053	0.012	-0.086
Population ages 40-44, female (% f)	-0.136	0.002	-0.132	0.025	-0.01	-0.008	-0.043
Population ages 45-49, female (% f)	-0.155	-0.016	-0.043	0.02	-0.001	-0.032	-0.012
Population ages 50-54, female (% f)	-0.158	-0.002	0.008	0.025	0.002	-0.037	0.01
Population ages 55-59, female (% f)	-0.157	0.017	0.038	0.063	0.027	-0.044	0.014
Population ages 60-64, female (% f)	-0.155	0.008	0.08	0.085	0.042	-0.045	0.012
Population ages 65 and above (%)	-0.148	0.011	0.129	0.064	0.081	-0.001	0.004
Population ages 65-69, female (% f) Population ages 70-74, female (% f)	-0.149 -0.146	$0.007 \\ 0.014$	$0.11 \\ 0.13$	$0.094 \\ 0.054$	$0.065 \\ 0.076$	-0.035 -0.004	$0.008 \\ 0.004$
Population ages 75-79, female (% f) Population ages 80 and above, female (%	-0.142 -0.14	$0.016 \\ 0.013$	$0.13 \\ 0.133$	$0.098 \\ 0.072$	$0.087 \\ 0.099$	-0.012 0.016	$0.012 \\ 0.001$
f) Population density (people per sq. km of	-0.027	0.014	-0.043	-0.06	-0.045	0.03	-0.092
land area)							
Population growth (annual %)	0.119	-0.028	-0.014	-0.065	0.087	0.055	-0.063
Population, female	-0.01	-0.251	0.005	0.034	-0.086	-0.011	0.008
PPP conversion factor, GDP (LCU per	0.024	-0.043	-0.082	0.063	0.037	-0.129	-0.049
international \$)							
Preprimary education, duration (years) Price level ratio of PPP conversion factor	-0.067 -0.093	-0.004 0.023	$0.003 \\ 0.076$	0.035 -0.209	$0.036 \\ 0.074$	$-0.085 \\ 0.071$	0.068 -0.054
(GDP) to market exchange rate							
Primary education, duration (years)	0.032	0.013	0.052	-0.277	-0.024	0.032	-0.018
Primary school starting age (years)	0.03	-0.018	0.049	0.12	0.095	-0.062	-0.028
Regulatory Quality: Estimate	-0.135	0.016	0.064	-0.106	0.025	0.031	-0.106
Renewable energy consumption (% of	0.107	-0.007	0.161	-0.002	0.02	-0.089	-0.042
total final energy consumption)							
Rule of Law: Estimate	-0.131	0.025	0.058	-0.152	0.028	0.033	-0.117
Rural population	0.003	-0.205	0.001	0.046	-0.108	-0.018	0.001
Rural population growth (annual $\%$)	0.103	-0.001	0.06	-0.005	0.048	0.001	-0.03
Scientific and technical journal articles	-0.049	-0.265	0.047	-0.03	0.023	0.071	-0.008
Secondary education, duration (years)	-0.026	0.007	0.024	0.225	0.052	-0.02	-0.055
Secure Internet servers	-0.036	-0.145	0.063	-0.067	0.083	0.113	0.003
Services, value added (% of GDP)	-0.108	0.044	0.057	-0.14	-0.1	0.116	-0.012
Sex ratio at birth (male births per female	-0.072	-0.068	-0.069	0.128	-0.041	-0.124	0.014
births)							
Surface area (sq. km)	-0.019	-0.184	-0.004	-0.008	0.078	0.016	0.033
Survival to age 65, female (% of cohort)	-0.149	0.005	-0.097	0.018	-0.013	0.043	0.034
Terrestrial and marine protected areas (% $$	-0.026	-0.019	0.089	-0.022	0.124	-0.055	-0.118
of total territorial area) Terrestrial protected areas (% of total	-0.033	0.021	0.075	-0.089	0.074	-0.093	-0.086
land area) Total fisheries production (metric tons)	-0.019	-0.241	-0.003	0.013	-0.067	-0.058	-0.006
Total greenhouse gas emissions excluding	0.034	-0.005	-0.071	-0.071	0.091	-0.02	-0.025
	0.054	-0.003	-0.071	-0.071	0.091	-0.02	-0.020
LULUCF (% change from 1990) Total greenhouse gas emissions excluding	-0.066	-0.013	-0.141	-0.034	0.133	0.084	-0.101
LULUCF per capita (t CO2e/capita) Total natural resources rents (% of GDP)	0.063	-0.016	-0.179	0.026	0.236	-0.01	-0.06

Table 3: PCA Loadings (continued)

variable	PC1	PC2	PC3	PC4	PC5	PC6	PC7
Tuberculosis case detection rate (%, all	-0.128	0.021	-0.031	-0.016	-0.045	0.053	0.049
forms) Urban population	-0.022	-0.275	0.008	0.017	-0.056	-0.001	0.013
Urban population growth (annual %) Voice and Accountability: Estimate	0.132 -0.107	-0.049 0.054	$0.007 \\ 0.144$	-0.041 -0.161	0.048 -0.005	0.014 -0.047	-0.058 -0.007

Table 4: Country Clusters and Status

Cluster	Developing	Developed
1	Afghanistan, Angola, Burundi, Benin,	
	Burkina Faso, Central African Republic,	
	Cote d'Ivoire, Cameroon, Congo, Dem. Rep.,	
	Congo, Rep., Comoros, Ethiopia, Gabon,	
	Ghana, Guinea, Gambia, The,	
	Guinea-Bissau, Equatorial Guinea, Kenya,	
	Liberia, Lesotho, Madagascar, Mali,	
	Mozambique, Mauritania, Malawi, Namibia,	
	Niger, Nigeria, Papua New Guinea, Rwanda,	
	Sudan, Senegal, Solomon Islands, Sierra	
	Leone, Somalia, Sao Tome and Principe,	
	Eswatini, Chad, Togo, Timor-Leste,	
	Tanzania, Uganda, Yemen, Rep., Zambia,	
	Zimbabwe	
2	Aruba, Antigua and Barbuda, Bahamas,	Australia, Bermuda, Canada, Switzerland,
	The, Barbados, Chile, Costa Rica, Dominica,	Cyprus, Denmark, United Kingdom,
	Grenada, Guam, Hong Kong SAR, China, St.	Greenland, Ireland, Iceland, Israel, Korea,
	Kitts and Nevis, St. Lucia, Macao SAR,	Rep., Luxembourg, Malta, Norway, New
	China, Mauritius, Palau, French Polynesia,	Zealand, Sweden
	Singapore, Seychelles, Thailand, Trinidad	
	and Tobago, Uruguay, St. Vincent and the	
	Grenadines	

3	Argentina, Bangladesh, Belize, Bolivia,	
	Brazil, Bhutan, Botswana, Colombia, Cabo	
	Verde, Dominican Republic, Ecuador, Egypt,	
	Arab Rep., Fiji, Micronesia, Fed. Sts.,	
	Guatemala, Guyana, Honduras, Haiti,	
	Indonesia, India, Jamaica, Jordan, Kyrgyz	
	Republic, Cambodia, Kiribati, Lao PDR,	
	Lebanon, Sri Lanka, Morocco, Mexico,	
	Marshall Islands, Myanmar, Malaysia,	
	Nicaragua, Nepal, Nauru, Pakistan, Panama,	
	Peru, Philippines, Paraguay, West Bank and	
	Gaza, El Salvador, Suriname, Syrian Arab	
	Republic, Tonga, Tunisia, Turkiye, Viet	
	Nam, Vanuatu, Samoa, South Africa	
4	China	United States
5	United Arab Emirates, Azerbaijan, Bahrain,	
	Brunei Darussalam, Algeria, Iran, Islamic	
	Rep., Iraq, Kazakhstan, Kuwait, Libya,	
	Maldives, Mongolia, Oman, Qatar, Saudi	
	Arabia, Tajikistan, Turkmenistan,	
	Uzbekistan	
6	Armenia, Cuba, Georgia	Albania, Austria, Belgium, Bulgaria, Bosnia
		and Herzegovina, Belarus, Czechia,
		Germany, Spain, Estonia, Finland, France,
		Greece, Croatia, Hungary, Italy, Japan,
		Lithuania, Latvia, Moldova, North
		Macedonia, Montenegro, Netherlands,
		Poland, Portugal, Romania, Russian
		Federation, Serbia, Slovak Republic,
		Slovenia, Ukraine
7	Djibouti	

Code for Clustering Algorithm

Function

```
#' Hierarchical Cluster Analysis (HCA)
  #' Performs agglomerative hierarchical clustering using either single or
     complete linkage.
4 # '
  #' @param D A square distance matrix (NxN) with pairwise distances between
     observations.
  #' @param linkage A string specifying the linkage method; either "single"
     or "complete".
7 #' @param check_matrix_validity A boolean indicating whether to check if D
     is a valid
             distance matrix (symmetric with diag 0 and non-negative off
     diags).
  # '
  #' Oreturn A list containing:
  #' \describe{
       \item{merge}{A matrix representing the sequence of merges, similar to
      'hclust' output.}
       \operatorname{height}_{A} numeric vector of the distances at which each merge
     occurred.}
       \item{merge_matrix}{A matrix showing cluster assignments at each step.}
  #'}
15
16
17 hca=function(D, linkage=c("single", "complete"),
     check_matrix_validity=TRUE){
    "Explanation see above"
    N=nrow(D) #Number of Observations stored for convenience
19
20
    if (check_matrix_validity){
21
      msg="Not a correctly specified distance matrix"
22
      #check if matrix is square
23
      if(nrow(D)!=ncol(D)){stop(msg)}
25
      #Check each element for symmetry, non-negativity and diags do be 0
      for (i in 1:(N)){
27
        for (j in 1:(N)){
```

```
if (i==j){if(D[i,j]!=0){stop(msg)}}
29
           else if (D[i,j]!=D[j,i] | D[i,j]<=0){stop(msg)}</pre>
30
         }}
31
       #print success, if matrix invalid, we would stop before here
32
       print("Matrix seems valid")}
33
    #Initialize lingage function:
35
     if (tolower(linkage) == "single") {
36
       linkage_calculator=function(g, h, clusters, D){
37
         distances=list()
39
         #extract index of observations in each cluster
         idx_g=which(clusters==g)
41
         idx_h=which(clusters==h)
43
         #get sub-matrix of D
         dist_matrix = D[idx_g, idx_h]
45
         #find minimum of distance
         min_dist_idx = which.min(dist_matrix)
47
         #convert the indices back to matrix index
49
         min_g = idx_g[(min_dist_idx - 1) %% length(idx_g) + 1]
50
         min_h = idx_h[(min_dist_idx - 1) %/% length(idx_g) + 1]
51
         return(list(min_distance = dist_matrix[min_dist_idx], min_g = min_g,
53
            min_h = min_h))
54
       }}
55
     else if (tolower(linkage) == "complete") {
56
       linkage_calculator=function(g, h, clusters, D){
57
         distances=list()
59
         #extract index of observations in each cluster
         idx_g=which(clusters==g)
61
         idx_h=which(clusters==h)
63
         #get sub-matrix of D
         dist_matrix = D[idx_g, idx_h]
65
         #find maximum of distance
         min_dist_idx = which.max(dist_matrix)
67
```

```
#convert the indices back to matrix index
69
         min_g = idx_g[(min_dist_idx - 1) %% length(idx_g) + 1]
70
         min_h = idx_h[(min_dist_idx - 1) %/% length(idx_g) + 1]
71
72
         return(list(min_distance = dist_matrix[min_dist_idx], min_g = min_g,
73
            min_h = min_h))
       }
74
     }
75
     else {stop("unimplemented linkage specification")}
76
     #Initialize numeric vector with one cluster per observation
78
     clusters=as.numeric(1:N)
79
80
     #get lists we will append to keep track of results
     merged_distance=numeric() #distances (height) for the merges
82
     merge_history=clusters # matrix of merges at each step
     h_list=numeric() #clusters that were merged into g
84
     g_list=numeric() #clusters that got merged into
85
86
     #initialize search list for all combinations, so that we don't have to
        get them each iteration
     #merged clusters will be removed successively
     search_combinations=as.matrix(t(combn(1:N, 2)))
89
90
91
     k=N #initialize iteration counter
93
94
     while(k>1){
95
96
97
       #update iteration counter
98
       k=k-1
99
100
       #initialize min distance and corresponding index:
101
       min_dist=Inf
102
       merge_g=NA
103
       merge_h=NA
104
105
       for (pair in 1:nrow(search_combinations)){
106
```

107

```
#calculate linkage
108
          linkage=linkage_calculator(search_combinations[pair,1],
109
             search_combinations[pair,2], clusters, D)
110
          #update minimum distance and pairs
111
          if (linkage$min_distance<min_dist){</pre>
112
            min_dist=unlist(linkage$min_distance)
113
            merge_g=search_combinations[pair,1]
114
            merge_h=search_combinations[pair,2]
115
         }
       }
117
118
       #update cluster vector
119
       clusters[clusters==merge_h]=merge_g
120
121
       #update merge_history matrix
122
       merge_history=cbind(merge_history, clusters)
123
       #update lists for merged clusters
125
       h_list=c(h_list, merge_h)
126
       g_list=c(g_list, merge_g)
127
129
       #append merge distance, checking that it is larger than all before
130
       if (length(merged_distance) == 0) {
131
          merged_distance=c(merged_distance, unlist(min_dist))
133
       else if (min_dist>max(merged_distance)){
134
         merged_distance=c(merged_distance, unlist(min_dist))
135
          }
136
       else{stop(paste("Minimum distance decreased, should not happen.
137
           K/N: ", k, "/", N))
138
       #Remove search options from combinations
139
       search_combinations <- search_combinations[search_combinations[,1] !=</pre>
140
           merge_h, , drop=FALSE]
       search_combinations <- search_combinations[search_combinations[,2] !=</pre>
141
           merge_h, , drop=FALSE]
     }
142
     #since we are always merging to the cluster with lower index, we can
143
        check if we correctly arrived at all cluster 1
```

```
if(any(as.integer(clusters))!=1){stop("Did not arrive at all in one
144
        cluster")}
145
     #---- finalize data outputs
146
147
     #get merge history matrix
148
     merge_history=(as.matrix(merge_history))
149
     colnames(merge_history) <-paste("Step",1:nrow(merge_history))</pre>
150
151
     #same output as merge in hclust package, just wiht different naming
152
        convention
     h_list=unname(h_list)
153
     g_list=unname(g_list)
154
     merge=as.matrix(t(rbind(g_list,h_list)))
156
     #same output as height in hclust package
157
     merged_distance=unname(merged_distance)
158
     return(list(merge=merge, height=merged_distance,
160
        merge_matrix=merge_history))
161 }
```

Testing Script

```
1 #Simple script simulating data an testing the hca function against hclust
 set.seed(1220)
source('/Users/johannesrenz/Library/Mobile
     Documents/com~apple~CloudDocs/BDS/ML2/Final_report/function/hca.R')
  gen_data=function(n){
     #normal blob cluster
     cluster1 <- data.frame(</pre>
       x = rnorm(n/2, mean = 0, sd = 1.5),
       y = rnorm(n/2, mean = 0, sd = 1.5),
       group = factor(1))
10
     #concentric circles
11
     angles \leftarrow runif(n/2, 0, 2 * pi)
12
     radii \leftarrow rep(10, n/2)
13
     circles <- data.frame(</pre>
14
       x = radii * cos(angles) + rnorm(n/2, sd = 0.3),
15
```

```
y = radii * sin(angles) + rnorm(n/2, sd = 0.3),
       group = factor(2))
17
    #Combine
18
     data <- rbind(cluster1, circles)</pre>
19
    return(data)
20
  }
21
22
  data=gen_data(100)
23
24
  #Dist object for package
  distances <- (dist(data[, c("x", "y")], method = "euclidean"))</pre>
26
27
  #Matrix input for my function
28
  distance_matrix <- as.matrix(distances)</pre>
30
  #Run for single linkage
32
  clusters=hca(distance_matrix, linkage="single")
  clusters_package=hclust(distances, method="single")
34
35
  if(all(clusters$height == clusters_package$height)){
36
     correct=1} else {print("heights dont match exactly")}
38
  #Run for single linkage
  clusters=hca(distance_matrix, linkage="complete")
  clusters_package=hclust(distances, method="complete")
42
  if(all(clusters$height == clusters_package$height)){
43
     if (correct==1){
44
       cat("\n\n", "Successful, all merges are done with the same distances
45
          using either linkage")}
  } else {print("heights dont match exactly")}
  Testing Script Output
  > source("~/Library/Mobile
     Documents/com~apple~CloudDocs/BDS/ML2/Final_report/function/hca_test.R",
      echo=F)
  [1] "Matrix seems valid"
```

```
Successful, all merges are done with the same distances using either
```

4 [1] "Matrix seems valid"

linkage

Code for the Report

```
#Analyse data
  rm(list=ls())
  library(rworldmap)
  library(NbClust)
  library(ggalt)
  library(dplyr)
  library(tidyr)
 library(ggplot2)
  library(xtable)
 library(readr)
  library(biplotEZ)
 library(FactoMineR)
  library(readxl)
  library(stargazer)
  library(Rfast)
  library(kableExtra)
  library(RColorBrewer)
  library(missMDA)
21
  #Set parameters:
23
  textsize=20
25
  #Set Parameter for Clustering and dimensional reduction
  set.seed(1220) #Since Cluster Starting is random, we set a seed
  K=7 # K is the number of clusters and sets the number of retained principle
     components
  D=K+1 #D is the number of principle components to retain
30
32
  #Utils:
34
  #Tex formatting
35
  escape_latex <- function(text) {</pre>
    text <- gsub("(% of total population)", "%", text)</pre>
37
    text <- gsub("female (% of female population)", "% f", text)</pre>
```

```
text <- gsub("(% of female population)", "% f", text)</pre>
     text <- gsub("\\\", "\\\textbackslash{}", text)</pre>
40
     text <- gsub("&", "\\\&", text)
41
     text <- gsub("%", "\\\%", text)
42
     text <- gsub("\\$", "\\\\$", text)
43
     text <- gsub("#", "\\\#", text)
44
     text <- gsub("_", "\\\\_", text)
45
     text <- gsub("\\{", "\\\\{", text)
46
     text <- gsub("\\}", "\\\\}", text)
47
     return(text)
  }
49
  #Number formatting
  format_number <- function(x) {</pre>
51
     x_rounded <- round(x, 4)
     if (abs(x_rounded) < 0.0001) {</pre>
53
       return("0") #if abs value is below 0.0001, display as "0"
     } else if (abs(x_rounded) >= 10^(-3) & abs(x_rounded) <= 10^3) {
55
       return(formatC(x_rounded, format = "f", digits = 4))
     } else {
57
       return(formatC(x_rounded, format = "e", digits = 4))
59
  }
61
62
  setwd('/Users/johannesrenz/Library/Mobile
      Documents/com~apple~CloudDocs/BDS/ML2/Final_report')
64
  df=read_csv("./Data/WB/2c50c149-0365-46a2-9545-ac89f31b79eb_Data.csv")
65
66
  #correctly specify NA
  df[df == ".."] <- NA
69
70
  #Rename Columns
  df <- df %>% rename(
     country = 'Country Name',
73
     alpha3 = 'Country Code',
74
     series = 'Series Name',
75
     var = '2018 [YR2018]'
77
78 print("----")
```

```
print(paste("Percentage of Missing Values before handling anything:",
                round(sum(is.na(df$var))/length(df$var),2)))
80
81
   #drop invalid obs
82
   df <- df %>%
83
     filter(!is.na(country))
85
   #Read UN Developing Nations Classification
   un_classification=read_excel('./Data/UN/classification_nations_UN.xlsx')
87
   developing_nations=un_classification%>%
     select(alpha3, status, m49)
89
90
   #Read Regions
91
   regions=read_excel("./Data/UN/regions.xlsx")
   #Filter subregions
   regions = regions[grepl("^[^0]*0$", regions$region_code), ]
   #merge regions and classification
95
   regions = merge(developing_nations, regions, by = "m49", all.x = TRUE)
   regions = regions%>%
     select(alpha3, region, status)
99
   #group regions into continents:
   regions $Continent <- ifelse (regions $region %in% c("Latin America and the
101
      Caribbean"), "Latin America, Caribbean",
                                 ifelse(regions$region %in% c("Northern
102
                                    America"), "North America",
                                        ifelse(regions$region %in% c("Southern
103
                                           Asia", "South-eastern Asia", "Eastern
                                           Asia"), "Asia",
                                                ifelse(regions$region %in%
104
                                                   c("Sub-Saharan Africa",
                                                   "Northern Africa"), "Africa",
                                                       ifelse(regions$region %in%
105
                                                          c("Southern Europe",
                                                          "Western Europe",
                                                          "Eastern Europe",
                                                          "Northern Europe"),
                                                          "Europe",
                                                              ifelse (regions $ region
106
                                                                 %in% c("Western
                                                                 Asia", "Central
```

```
ifelse(regions$reg;
107
                                                                               "Oceania",
                                                                               "Oceania",
                                                                               NA))))))
108
109
110
111
112
   df <- merge(df, regions, by = "alpha3")</pre>
113
114
   #transform data to wide format
   df <- df %>%
116
     select(country, region, Continent, alpha3, series, status, var)%>%
117
     pivot_wider(names_from = series, values_from = var)
118
120
   #Dropping data
121
122
   df = as.data.frame(df)
   df <- df[, sort(names(df))]</pre>
124
125
   #Identify the first occurrence of each base name
126
   col_base_names <- sub("[,|:].*", "", colnames(df))</pre>
   col_base_names <- sub("\\(.*", "", col_base_names)</pre>
128
   col_base_names <- trimws(col_base_names)</pre>
129
130
   #keep only one measure per indicator
131
   df <- df[, !duplicated(col_base_names)]</pre>
132
133
   #initially drop vars with very low observations
134
   df <- df[, colSums(is.na(df)) <= 0.1*nrow(df)]#drop low obs obs</pre>
135
   df \leftarrow df[rowSums(is.na(df)) \leftarrow 0.1*ncol(df),]#drop low obs vars
136
137
   rownames (df) = df $ country
138
139
   #split into labels and data
   df_lab=df%>%select(country, region, Continent, alpha3, status)
   df=df%>%select(-country, -region, -Continent, -alpha3, -status)
```

Asia"), "Asia",

```
143
   #Convert type to num
144
   df[] <- lapply(df, function(x) as.numeric(as.character(x)))</pre>
145
146
   #Some diagnostic Printing
147
   cat("\n -----\nMissing Countries are:\n",
148
       paste(un_classification$Country[un_classification$alpha3 %in%
149
          setdiff(un_classification$alpha3, df_lab$alpha3)] , collapse ="\n"))
150
   cat("\n\n -----\nFraction of NA to be imputed is:",
151
       sum(is.na(df))/sum(is.na(df),!is.na(df)),
152
       "\n")
153
154
   cat("\n\nDistribution of NA across Countries:\n")
   print(summary(colSums(is.na(df))))
156
   print(quantile(colSums(is.na(df)), probs = seq(0.1, 1, by = 0.05)),
      digits=1)
   cat("\n\nDistribution of NA across Variables:\n")
159
   print(summary(rowSums(is.na(df))))
160
   print(quantile(rowSums(is.na(df)), probs = seq(0.1, 1, by = 0.05)),
161
      digits=1)
162
   #Plot distribution of missings
163
   data_variable <- data.frame(Missing = colSums(is.na(df)), Type = "Variable")</pre>
164
   data_country <- data.frame(Missing = rowSums(is.na(df)), Type = "Country")</pre>
165
166
   ggplot(data_variable, aes(x = Missing)) +
167
     geom_histogram(binwidth = 1, color = "black", fill = "grey") +
168
     labs(x = "Missing Values", y = "Frequency") +
169
     theme_minimal()+
170
     theme (
171
       text = element_text(size = textsize),
172
       axis.title = element_text(size = textsize),
173
       axis.text.x = element_text(size = textsize),
174
       axis.text.y = element_text(size = textsize)
175
176
   ggsave("./figures/missing_var.png", width = 8, height = 6)
177
178
179
   ggplot(data_country, aes(x = Missing)) +
```

```
geom_histogram(binwidth = 1, color = "black", fill = "grey") +
181
     labs(x = "Missing Values", y = "Frequency") +
182
     theme_minimal()+
183
     theme (
184
        text = element_text(size = textsize),
185
        axis.title = element_text(size = textsize),
186
        axis.text.x = element_text(size = textsize),
187
        axis.text.y = element_text(size = textsize)
188
189
   ggsave("./figures/missing_country.png", width = 8, height = 6)
191
192
   cat("\n -----\n")
193
   df = as.data.frame(df)
195
   #imput NA using PCA
197
   ndim=estim_ncpPCA(df, scale=TRUE) # estimate optimal number of PCA
      dimensions to use in imputation
   impPCA=imputePCA(df, ncp=ndim$ncp, nboot=1, scale=TRUE) #for bootstrap
      replace with MIPCA
200
   #When using MIPCA
201
   #plot(impPCA, choice="dim")
202
   #this gives a feeling for the uncertainty of the components. did not fit in
203
      the paper due to length contraints
204
205
206
   tmp = df
207
   df = as . data . frame (impPCA $ completeObs)
208
209
   rownames(df)=rownames(tmp)
210
   colnames(df) = colnames(tmp)
211
212
   #--- Summary Statistics
213
214
   #function to create a tex table with kableExtra
215
   create_latex_summary_table <- function(data, file_name) {</pre>
     summary_stats <- data %>%
217
        summarise(across(everything(), list(
218
```

```
Mean = ~ mean(.x, na.rm = TRUE),
219
          SD =   ^{\sim} sd(.x, na.rm = TRUE), 
220
          Min =  min(.x, na.rm = TRUE),
221
          Max = ^max(.x, na.rm = TRUE)
222
        ))) %>%
223
        pivot_longer(everything(), names_to = c("Variable", ".value"),
224
           names_sep = "_")
225
     summary_stats$Variable <- gsub("_", "_\\\newline ",</pre>
226
         summary_stats$Variable)
     summary_stats$Variable <- sapply(summary_stats$Variable, escape_latex)</pre>
227
228
     summary_stats[, 2:5] <- lapply(summary_stats[, 2:5], function(col)</pre>
229
         sapply(col, format_number))
230
     table_latex <- summary_stats %>%
231
       kbl(format = "latex", booktabs = TRUE, longtable = TRUE, escape =
232
           FALSE, align = "lcccc",
            caption = "Descriptive Statistics Summary", label = "tab:app:sum")
233
        kable_styling(latex_options = c("repeat_header"), font_size = 9) %>%
234
        column_spec(1, width = "8cm")
235
     writeLines(table_latex, file_name)
236
   }
237
   #run func
238
   create_latex_summary_table(df, "./tables/summary_appendix.tex")
239
240
241
   write_csv(df, "./Data/data_clean.csv")
242
   write_csv(df_lab, "./Data/data_clean_labels.csv")
243
244
245
246
247
   #Data Analysis Section
249
250
251
   setwd('/Users/johannesrenz/Library/Mobile
252
      Documents/com~apple~CloudDocs/BDS/ML2/Final_report')
   df=read_csv("./Data/data_clean.csv")
```

```
df_lab=read_csv("./Data/data_clean_labels.csv")
255
  #---- ---- ----
256
  #----Analysis Part
257
  #---- ---- ----
258
259
  #Convert df to Z-scores
260
  df=scale(df, center = TRUE, scale = TRUE)
261
262
263
  #Run the PCA using SVD
264
  SVD=svd(df)
265
266
  var_expl=SVD$d^2 / sum(SVD$d^2)
   cumu_var_expl = cumsum(var_expl)
268
269
  #manually compute the transformed coordinates by multiplying the centered
270
     data by PCs
  #gives the projections of the data onto the principal components
   transformed_coordinates = df %*% SVD$v
273
  #limti to the first k transformed coordinates (dimensions 1 to number of
     clusters -1)
  transformed_coordinates_k = transformed_coordinates[, 1:D]
276
  #perform k-means clustering on the first k components
  kmeans_result = kmeans(transformed_coordinates_k,
278
                        centers = K,
279
                        nstart=10000)
280
281
282
  #---- ---- ----
283
  #Results section
284
  #---- ---- ----
285
  #plotting and tables from here onwards
  #---- ---- ----
287
  loadings=data.frame(variable=colnames(df),loading=SVD$v[,1])
289
   loadings$loading=round(loadings$loading, digits=4)
291
292
```

```
293
   #df with PCA component names
294
   pc_df <- as.data.frame(round(SVD$v[, 1:D], 3))</pre>
295
   colnames(pc_df) <- paste0("PC", 1:D)</pre>
296
   pca_print <- data.frame(variable = colnames(df), pc_df)</pre>
297
298
   #PCA long table with cumulative variance explained
299
   create_PCA_long_table <- function(data, cumu_var_expl, file_name) {</pre>
300
     data_selected <- data %>% mutate(across(everything(), escape_latex))
301
     data_selected <- data_selected %>% mutate(across(where(is.numeric),
302
        round, digits = 2))
303
     #Create header for cumulative variance explained
304
     header_variance <- c("", paste0("(", round(cumu_var_expl[1:(ncol(data)-1)]
305
        * 100, 2), "%)"))
     header_combined <- c(header_pc, header_variance)
306
307
     table_latex <- data_selected %>%
308
       kbl(format = "latex", booktabs = TRUE, longtable = TRUE, escape = FALSE,
309
            align = "lcccc", caption = "PCA Loadings", label = "tab:app:pca")
310
       kable_styling(latex_options = c("repeat_header"), font_size = 9) %>%
       column_spec(1, width = "8cm") %>%
312
       add_header_above(header_combined)
313
314
     write_lines(table_latex, file_name)
316
317
318
   create_PCA_long_table(pca_print, cumu_var_expl,
319
      "./tables/loadings_appendix.tex")
320
321
322
323
   #print the cumulative variance explained
324
   cat("\n Cumulative Variance Explained by
325
      Component:",round(cumu_var_expl[1:10],2))
   cat("\n Using our set of dimensions, we
      explain:",round(cumu_var_expl[D],2), "% of variance")
327
```

```
328
   #df to make plotting easier
329
   df_plot <- data.frame(</pre>
330
     X = transformed_coordinates[, 1],
331
     Y = transformed_coordinates[, 2],
332
     Region = df_lab$region,
333
      Country = df_lab$country,
334
      alpha3 = df_lab$alpha3,
335
     Continent = df_lab $ Continent,
336
     Status=df_lab$status,
337
      Cluster = as.factor(unname(kmeans_result$cluster)))
338
339
340
   #map plot
   world_map <- getMap()</pre>
342
   map_data <- merge(world_map, df_plot, by.x = "ISO_A3", by.y = "alpha3",</pre>
       all.x = TRUE)
   colors <- brewer.pal(length(unique(df_plot$Cluster)), "Set3")</pre>
       colors in 'Set3'
345
   map_data$dev=ifelse(map_data$Status=="Developed", 1,0.4)
346
   png("./figures/world_map.png")
347
   mapCountryData(map_data, nameColumnToPlot = "Cluster",
348
                    mapTitle = "",
349
                    colourPalette = colors,
350
                    catMethod = 'categorical',
351
                    borderCol="#808080",
352
                    nameColumnToHatch='dev',
353
                    addLegend =
                                   T)
354
   dev.off()
355
356
357
358
   print(table(df_plot$Region, df_plot$Cluster, df_plot$Status))
359
360
   table_data <- table(df_plot$Cluster, df_plot$Status)
361
   print(table_data)
362
   for (cluster in 1:nrow(table_data)) {
363
     developed_fraction <- table_data[cluster, "Developed"] /</pre>
364
         sum(table_data[cluster, ])
     minority_group <- ifelse(table_data[cluster, "Developed"] <</pre>
365
```

```
table_data[cluster, "Developing"], "Developed", "Developing")
     cat("\n Cluster", cluster, "- Developed Fraction:",
366
        round(developed_fraction,2), "%.\n Minority group:", minority_group,
        "\n Countries in the minority group:\n")
     print(df_plot[df_plot$Cluster == cluster & df_plot$Status ==
367
        minority_group, "Country"])
   }
368
369
   table_df <- data.frame(
370
     Cluster = 1:nrow(table_data),
371
     Developed = table_data[, "Developed"],
372
     Developing = table_data[, "Developing"],
373
     Fraction = round(table_data[, "Developed"] / rowSums(table_data) * 100,
374
        1),
     Minority = ifelse(table_data[, "Developed"] < table_data[, "Developing"],
375
        "Developed", "Developing"),
     Minority_Countries = sapply(1:nrow(table_data), function(cluster) {
376
       countries <- df_plot[df_plot$Cluster == cluster & df_plot$Status ==</pre>
377
           ifelse(table_data[cluster, "Developed"] < table_data[cluster,</pre>
           "Developing"], "Developed", "Developing"), "Country"]
       if (length(countries) > 0) paste(countries, collapse = ", ") else "None"
378
     })
   )
380
381
   tex=kbl(table_df, format = "latex", booktabs = TRUE,
382
       caption = "Cluster Composition and Minority Groups", label = "cluster")
383
          %>%
     column_spec(6, width = "7cm") %>% # Use p{} for proper text wrapping
384
     footnote(general = "Note: The Developed Fraction represents the
385
        percentage of developed countries in each cluster.")
   write_lines(tex, "./tables/cluster_assignment.tex")
386
387
388
389
390
391
   #cluster compactness plot
   nc <- NbClust(data=transformed_coordinates_k, min.nc = 2, max.nc = 25,</pre>
393
                  method = "kmeans", index = "tracew")
394
395
   index_data <- data.frame(Index = 2:25, Cluster = nc$All.index,</pre>
```

```
var_expl=var_expl[1:24], cumulative_var_expl=cumu_var_expl[1:24])
397
398
   p = ggplot(index_data, aes(x = Index)) +
399
     #plot variance explained as a line
400
     geom_line(aes(y = var_expl), color = "blue") +
401
     geom_point(aes(y = var_expl), color = "blue") +
402
     #plot cumulative variance explained as a second line
403
     geom_line(aes(y = cumulative_var_expl * max(index_data$var_expl) /
404
        max(index_data$cumulative_var_expl)), color = "red") +
     geom_point(aes(y = cumulative_var_expl * max(index_data$var_expl) /
405
        max(index_data$cumulative_var_expl)), color = "red") +
     theme_minimal() +
406
     labs(
407
       x = "Principal Component",
408
       y = "Variance Explained"
410
     #secondary axis scaled so max is 1 but data remains unchanged
     scale_y_continuous(
412
       sec.axis = sec_axis(
413
          ~ . * max(index_data$cumulative_var_expl) / max(index_data$var_expl),
414
              # Adjust only axis scaling
         name = "Cumulative Variance Explained (Max = 1)"
415
       )
416
     ) +
417
     theme (
418
       text = element_text(size = textsize),
419
       axis.title = element_text(size = textsize),
420
       axis.text.x = element_text(size = textsize),
421
       axis.text.y = element_text(size = textsize)
422
     )
423
424
   print(p)
425
426
   ggsave("./figures/pca_scree.png", width = 8, height = 6)
428
429
   q = ggplot(index_data, aes(x = Index)) +
430
     geom_line(aes(y = Cluster), color = "black") +
431
     geom_point(aes(y = Cluster), color = "black") +
432
     theme_minimal() +
433
```

```
labs(
434
       x = "Number of Clusters",
435
       y = "Compactness"
436
     ) +
437
     theme(
438
       text = element_text(size = textsize),
439
       axis.title = element_text(size = textsize),
440
       axis.text.x = element_text(size = textsize),
441
       axis.text.y = element_text(size = textsize)
442
     )
443
444
   print(q)
445
446
   ggsave("./figures/cluster_scree.png", width = 8, height = 6)
448
   #Appendixx table version of map
   df_latex <- df_plot %>%
450
     group_by(Cluster, Status) %>%
451
     summarise(Countries = paste(Country, collapse = ", "), .groups = "drop")
452
        %>%
     pivot_wider(names_from = Status, values_from = Countries, values_fill =
453
        list(Countries = ""))
454
   groups_clusters_table <- kable(df_latex, format = "latex", booktabs = TRUE,
455
                          longtable = TRUE, escape = FALSE,
456
                          caption = "Country Clusters and Status",
457
                          label = "map_table") %>%
458
     column_spec(2, width = "8cm") %>%
459
     column_spec(3, width = "8cm")
460
461
462
   write_lines(groups_clusters_table, "./tables/map_table.tex")
```