

# Clustering / World 2023

Github Bavla OpenAlex [https://github.com/bavla/OpenAlex/blob/main/Countries/pics/world.md]

C:/Users/vlado/work/OpenAlex/API/world.R

#### Clustering of the Balassa index matrix

```
> wdir <- "C:/Users/vlado/work/OpenAlex/API"</pre>
> setwd(wdir)
> library(httr)
> library(jsonlite)
> source("https://raw.githubusercontent.com/bavla/Rnet/master/R/Pajek.R")
  source("OpenAlex4.R")
> library(gplots)
> CorEu <- function(W,p=1){
     sqrt(sum((W[u,]-W[v,])**2) -
(W[u,u]-W[v,u])**2 - (W[u,v]-W[v,v])**2 +
        p*((W[u,u]-W[v,v])**2 + (W[u,v]-W[v,u])**2))
     return(D)
> load("./years/Matrix2023.Rdata")
> P <- M; diag(P) <- 0
> Y <- as.character(2023); i <- 1
> D <- rowSums(P); T <- sum(D); n <- nrow(P)
> for(u in 1:(n-1)) for(v in (u+1):n) P[u,v] \leftarrow P[v,u] \leftarrow P[u,v]*T/D[u]/D[v]
> X <- Z <- log2(P)
> Z[Z == -Inf] <- 0; Z[is.nan(Z)] <- 0
> X[X == -Inf] <- NA; X[is.nan(X)] <- 0
> X[,D==0] <- NA; X[D==0,] <- NA</pre>
> t <- hclust(as.dist(CorEu(Z)),method="ward.D")</pre>
> plot(t,hang=-1,main="World 2023 / CorrEuclid / Ward",cex=0.3)
> pdf(file=paste("WorldBalassa",Y[i],".pdf",sep=""),width=30,height=30)
> heatmap.2(X,Rowv=as.dendrogram(t),Colv="Rowv",dendrogram="column",
  scale="none",revC=TRUE,col=bluered(100),na.color="yellow",
trace="none",density.info="none",keysize = 0.8,
   main=paste("World ",Y[i]," / Balassa / Ward",sep=""))
> dev.off()
```

Picture of the clustering of the Balassa index matrix [https://github.com/bavla/OpenAlex/blob/main/Countries/pics/WorldBalassa2023.pdf].

Picture of the corresponding intensities of co-authorship [https://github.com/bavla/OpenAlex/blob/main/Countries/pics/WorldCoA2023.pdf].

## Clustering of coauthorship intensities

```
> P <- M; diag(P) <- 0
> X <- Z <- log2(P)
> Z[Z == -Inf] <- 0; Z[is.nan(Z)] <- 0
> X[X == -Inf] <- Na; X[is.nan(X)] <- 0
> X[,D==0] <- Na; X[D==0,] <- Na
> h <- hclust(as.dist(CorEu(Z)),method="ward.D")
> myPalette <- colorRampPalette(c("white","black"))(n=100)
> pdf(file=paste("WorldCoA",Y[i],".pdf",sep=""),width=30,height=30)
> heatmap.2(X,Rowv=as.dendrogram(t),Colv="Rowv",dendrogram="column",
+ scale="none",revC=TRUE,col=myPalette,na.color="yellow",
+ trace="none",density.info="none",keysize = 0.8,
+ main=paste("WorldCoW",Y[i]," / log2 / Ward",sep=""))
> dev.off()
> pdf(file=paste("WorldCoW",Y[i],".pdf",sep=""),width=30,height=30)
> heatmap.2(X,Rowv=as.dendrogram(h),Colv="Rowv",dendrogram="column",
+ scale="none",revC=TRUE,col=myPalette,na.color="yellow",
+ trace="none",density.info="none",keysize = 0.8,
+ main=paste("World ",Y[i]," / log2 w / Ward",sep=""))
> dev.off()
```

Picture of the clustering of intensities [https://github.com/bavla/OpenAlex/blob/main/Countries/pics/WorldCoW2023.pdf].

#### Reordering clusters

A better picture can be obtained by flipping some subtrees in the dendrogram (see Reorder [https://github.com/bavla/NormNet/blob/main/data/natalija/reorder.md])

```
4. 8. 24, 07:22
   [5,] 244 235
   [6,] 245 242
    [7,] 246 232
                   244
                   238
   [8,] 247 237
   [9,] 248 233 245
  [10,] 249 246 247
  [11,] 250 248 249
  > h$merge <- flip(186,flip(237,flip(247,h$merge)))
  > pdf(file=paste("WorldCoT",Y[i],".pdf",sep=""),width=30,height=30)
  > heatmap.2(X,Rowv=as.dendrogram(h),Colv="Rowv",dendrogram="column",
  + scale="none",revC=TRUE,col=myPalette,na.color="yellow",
trace="none",density.info="none",keysize = 0.8,
      main=paste("World ",Y[i]," / log2 w / Ward",sep=""))
  > dev.off()
  > cbind(235:nm,h$merge[235:nm,])
        [,1] [,2] [,3]
   [1,] 235 178 183
[2,] 236 218 227
   [3,] 237 186
                    19
   [4,] 238 222
   [5,] 239 216
                   236
   [6,] 240 192 220
   [7,] 241 229
                   230
   [8,] 242 231 239
   [9,] 243 204
                   240
  [10,] 244 235
                   241
  [11,] 245 242
  [12,] 246 232
  [13,] 247
              238
                   237
  [14,] 248 233
                   245
  [15,] 249 246
                   247
```

Picture of the reordered intensity matrix [https://github.com/bavla/OpenAlex/blob/main/Countries/pics/WorldCoT2023.pdf].

### Reordering Balassa

[16,] 250 248 249

```
> toFather <- function(tm){</pre>
   n \leftarrow nrow(tm); T \leftarrow rep(0,2*n+1)
    for(i in 1:n){
      for(j in 1:2){
        p <- tm[i,j]
        if(p<0) T[-p] <- i+n+1 else T[n+1+p] <- i+n+1
    return(T)
+ }
> minCl <- function(u,v,T){</pre>
  if(min(u,v)==0) return(T[max(u,v)])
# cat(u," ",v,":",T[u]," ",T[v],"\n")
  if(u==v) return(u)
   return( if(T[u] < T[v]) minCl(T[u], v, T) else minCl(u, T[v], T) )
+ }
> F <- toFather(t$merge)
> N <- rownames(M)
> which(N=="SI")
[1] 205
> minCl(which(N=="AN"), which(N=="FR"), F) - n
[1] 239
> minCl(which(N=="IM"), which(N=="FR"), F) - n
[1] 230
> minCl(which(N=="IM"),which(N=="AW"),F) - n
[1] 220
> minCl(which(N=="ST"), which(N=="FR"), F) - n
[1] 225
> minCl(which(N=="TV"), which(N=="FK"), F) - n
[1] 148
> minCl(which(N=="PN"),which(N=="FK"),F) - n
[1] 129
> minCl(which(N=="VG"),which(N=="FK"),F) - n
[1] 78
> minCl(which(N=="SH"), which(N=="FK"), F) - n
[1] 72
> t$merge <- flip(225,flip(220,flip(230,flip(239,flip(250,t$merge)))))
> t$merge <- flip(24,flip(72,flip(78,flip(129,flip(148,t$merge)))))</pre>
> pdf(file=paste("WorldBalassaF",Y[i],".pdf",sep=""),width=30,height=30)
> heatmap.2(X,Rowv=as.dendrogram(t),Colv="Rowv",dendrogram="column",
   scale="none",revC=TRUE,col=bluered(100),na.color="yellow",
trace="none",density.info="none",keysize = 0.8,
   main=paste("World ",Y[i]," / Balassa / Ward",sep=""))
> dev.off()
```

Picture of the reordered Balassa matrix [https://github.com/bavla/OpenAlex/blob/main/Countries/pics/WorldBalassaF2023.pdf].

Europe

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