

## Statistical Significance of Dissimilarity Matrix using MRPP and ANOSIM

```
In [164]: library(vegan)
library(mefa)
```

```
In [170]: rankdata<-read.csv(file="/Users/Data/simple_data_rankings.csv")
rankdata <- rankdata[-1]
```

```
In [138]: rawdata<-read.csv(file="/Users/Data/simple_rawdata.csv")
rawdata <- rawdata[-1]
rawdata <- rawdata[-1,]
```

```
In [175]: for (j in 2:10){
  for (i in 1:(j-1)){
    rawdata[i,j] <- 0
    rankdata[i,j] <- 0
  }}

```

```
In [181]: #create distance matrix for raw data

col1 <- rawdata[1:10,1]
col2 <- rawdata[2:10,2]
col3 <- rawdata[3:10,3]
col4 <- rawdata[4:10,4]
col5 <- rawdata[5:10,5]
col6 <- rawdata[6:10,6]
col7 <- rawdata[7:10,7]
col8 <- rawdata[8:10,8]
col9 <- rawdata[9:10,9]
col10 <- rawdata[10:10,10]
vraw <- c(col1,col2,col3,col4,col5,col6,col7,col8,col9,col10)
raw.distance.matrix <- vec2dist(vraw,11)
```

```
In [182]: #create distance matrix for ranked data

col1 <- rankdata[1:10,1]
col2 <- rankdata[2:10,2]
col3 <- rankdata[3:10,3]
col4 <- rankdata[4:10,4]
col5 <- rankdata[5:10,5]
col6 <- rankdata[6:10,6]
col7 <- rankdata[7:10,7]
col8 <- rankdata[8:10,8]
col9 <- rankdata[9:10,9]
col10 <- rankdata[10:10,10]
vrnk <- c(col1,col2,col3,col4,col5,col6,col7,col8,col9,col10)
rank.distance.matrix <- vec2dist(vrnk,11)
```

```
In [189]: #create grouping of data
```

```
rdata <- factor(c(1,1,1,1,2,2,2,2,2,2,2),labels=c("Year 1","Year 2"))
```

```
In [200]: model.mrpp <- mrpp(raw.distance.matrix, rdata,permutations=10000)
```

```
In [198]: model.anosim <- anosim(rank.distance.matrix, rdata,permutations=10000)
```

```
In [202]: model.mrpp
```

Call:

```
mrpp(dat = raw.distance.matrix, grouping = rdata, permutations = 10000)
```

Dissimilarity index:

Weights for groups: n

Class means and counts:

	Year 1	Year 2
delta	0.1361	0.172
n	4	7

Chance corrected within-group agreement A: 0.1732

Based on observed delta 0.159 and expected delta 0.1923

Significance of delta: 0.0024998

Permutation: free

Number of permutations: 10000

```
In [210]: summary(model.anosim)
```

Call:

```
anosim(dat = rank.distance.matrix, grouping = rdata, permutations = 10000)
```

Dissimilarity:

ANOSIM statistic R: 0.5331

Significance: 0.0025997

Permutation: free

Number of permutations: 10000

Upper quantiles of permutations (null model):

	90%	95%	97.5%	99%
	0.212	0.280	0.380	0.454

Dissimilarity ranks between and within classes:

	0%	25%	50%	75%	100%	N
Between	14	25.75	35.5	45.25	55	28
Year 1	2	4.75	11.0	12.75	24	6
Year 2	1	8.00	21.0	37.00	52	21

```
In [207]: plot(model.anosim)
```

Warning message in bxp(structure(list(stats = structure(c(14, 25.5, 35.5, 45.5, :  
"some notches went outside hinges ('box'): maybe set notch=FALSE"

