

Multi-view Banded Spectral Clustering (mvBSC)

```
# devtools::install_github("celehs/mvBSC")
```

```
library(mvBSC)
```

```
library(data.table)
```

```
va_cosK <- readRDS(paste0("data/va_I00-I25_cosineMat.rds"))
bio_cosK <- readRDS(paste0("data/biobank_I00-I25_cosineMat.rds"))
this.R <- readRDS(paste0("data/I00-I25_distR_wt_avg_1.rds"))
icd.info <- readRDS("data/rollable_new_icd_info_20190130.rds")
codes_in_use <- colnames(bio_cosK)
codes_with_phecode <- codes_in_use[!is.na(icd.info[codes_in_use]$PheCode)]
pheCodes <- icd.info[codes_with_phecode]$PheCode
Z0 <- get_Z(codes_with_phecode, pheCodes)
delta0 <- min(apply(this.R, 1, max)) / 2
delta0
```

```
## [1] 24.5
```

```
initial <- mvbsc_fit(
  codes = rownames(this.R),
  distance = this.R,
  similarity = list(va_cosK, bio_cosK),
  ncluster = 30,
  weights = c(va.wt = 0.5, bio.wt = 0.5),
  delta = delta0,
  band = 4,
  seed = 123)
initial
```

```
## $cluster
```

```
##   394.9   395.9   396.0   396.1   396.2   396.3   396.8   396.9   397.0
##      28      28      25      25      25      25       4       4      28
##   397.1   397.9   398.99  401.0   401.1   401.9   402.00  402.01  402.10
##       4      25       4      10      25      10      21      10      21
##  402.11  402.90  402.91  403.00  403.01  403.10  403.11  403.90  403.91
##      21      21      10       9       9      14      14      14      10
##  404.00  404.01  404.02  404.03  404.10  404.11  404.12  404.13  404.90
##       9       9       9       9       9      14       9      30      14
##  404.91  404.92  404.93  405.01  405.09  405.11  405.19  405.91  405.99
##      14       9       9      29      29      29      25      29      29
##  410.00  410.01  410.02  410.10  410.11  410.12  410.20  410.21  410.22
##       3       3       3       3       3       3      19      16      16
##  410.30  410.31  410.32  410.40  410.41  410.42  410.50  410.51  410.52
##      16      16      16       2       2       2      19      19       2
##  410.60  410.61  410.62  410.70  410.71  410.72  410.80  410.81  410.82
##      19      16       2      19       5       5       3       2       5
##  410.90  410.91  410.92  411.1   411.89  413.0   413.9   414.00  414.01
##       2       2       2       5      26       5       5      23      23
##  414.02  414.03  414.04  414.05  414.06  414.07  414.10  414.19  414.8
##      26      26      26      26      23      23      26      26      26
##   414.9   429.2   429.79   I00.   I01.0   I01.1   I01.2   I01.8   I01.9
##      23      11       7      25      17      17      17      17      17
```

```

##      I02.0      I02.9      I05.0      I05.1      I05.2      I05.8      I05.9      I06.0      I06.1
##      25        25          1          1          1          15         15          1          28
##      I06.2      I06.8      I06.9      I07.1      I07.2      I07.8      I07.9      I08.0      I08.1
##      1          15         15         27         27         27         27         27         27
##      I08.2      I08.3      I08.8      I08.9      I09.1      I09.2      I09.81      I09.89      I09.9
##      27         27         27         27         25         25          4          4          4
##      I10.       I11.0      I11.9      I12.0      I12.9      I13.0      I13.10      I13.11      I13.2
##      25         25         25         30         18         18         18         30         30
##      I15.0      I15.1      I15.2      I15.8      I15.9      I20.0      I20.1      I20.8      I20.9
##      24         24         24         24         24          8          5          8          8
##      I21.01     I21.02     I21.09     I21.11     I21.19     I21.21     I21.29     I21.3      I21.4
##      20          8          8          20         20          8          8          8          8
##      I22.0      I22.1      I22.2      I22.8      I22.9      I23.1      I23.2      I23.6      I23.7
##      20         20          8          7          7          7          7          7          7
##      I23.8      I24.0      I24.1      I24.8      I24.9      I25.10     I25.110     I25.111     I25.118
##      7          7          7          6         12          6         13          6          6
##      I25.119     I25.2      I25.3      I25.41     I25.42     I25.5      I25.6      I25.700     I25.701
##      6          6         12         11         12          6         12         13         22
##      I25.708     I25.709     I25.710     I25.711     I25.718     I25.719     I25.720     I25.721     I25.728
##      11         13         13         22         13         22         13         22         22
##      I25.729     I25.739     I25.750     I25.758     I25.759     I25.760     I25.769     I25.790     I25.791
##      13         22         22         11         22         13         22         13         13
##      I25.798     I25.799     I25.810     I25.811     I25.812     I25.82      I25.83      I25.84      I25.89
##      22         22         13         11         25         13          6          6          6
##      I25.9
##      6
##
## $cluster_info
##      cluster size max_dist
## 1          22  11    0.09
## 2          16   6    0.28
## 3           2   9    0.31
## 4           3   7    0.40
## 5          19   5    0.51
## 6          18   3    1.40
## 7          11   5    1.41
## 8          13  12    1.41
## 9          23   5    1.60
## 10         29   5    1.60
## 11         17   5    1.80
## 12         21   4    1.80
## 13         24   5    1.80
## 14         12   4    2.40
## 15         26   8    3.01
## 16         20   5    3.19
## 17          4   7    3.20
## 18          6  11    3.20
## 19         14   6    3.20
## 20         15   4    3.20
## 21          1   5    3.40
## 22          9  10    3.40
## 23         30   4    3.40
## 24          5   7    3.80
## 25          7  10    4.60

```

```
## 26      27    10    4.60
## 27      28     4    5.13
## 28      10     5    6.00
## 29       8    10    6.40
## 30      25    16   96.61
```

```
cluster0 <- subset(initial$cluster_info, max_dist > delta0)$cluster
cluster0
```

```
## [1] 25
```

```
regroup <- vector("list", length(cluster0))
names(regroup) <- paste0("initial_", cluster0)
```

```
for (i in 1:length(cluster0)) {
  tmp <- names(initial$cluster[initial$cluster == cluster0[i]])
  for (k in 2:(length(tmp) - 1)) {
    try <- mvbsc_fit(
      codes = tmp,
      distance = this.R,
      similarity = list(va_cosK, bio_cosK),
      ncluster = k,
      weights = c(va.wt = 0.5, bio.wt = 0.5),
      delta = delta0,
      band = 4,
      seed = 123)
    if (all(try$cluster_info$max_dist <= delta0)) break
  }
  regroup[[i]] <- try
}
regroup
```

```
## $initial_25
## $initial_25$cluster
##   396.0  396.1  396.2  396.3  397.9  401.1  405.19  I00.  I02.0
##      6      6      6      6      6      3      7      8      6
##   I02.9  I09.1  I09.2  I10.  I11.0  I11.9 I25.812
##     10      1      2      5      4      4      9
##
## $initial_25$cluster_info
##   cluster size max_dist
## 1      1     1     0.0
## 2      2     1     0.0
## 3      3     1     0.0
## 4      5     1     0.0
## 5      7     1     0.0
## 6      8     1     0.0
## 7      9     1     0.0
## 8     10     1     0.0
## 9      4     2     1.8
## 10     6     6    21.2
```