Describing Bilteral Migration Data

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- Rogers et al. (2002) proposed dis-aggregating origin-destination flow tables into separate components to allow for an easier examination of migration flows
 - \bullet Overall component level of migration γ
 - ullet Origin component relative 'pushes' from each region $lpha_i$
 - Destination component relative 'pulls' to each region β_j
 - Origin–Destination interaction component physical or social distance between places not explained by the overall and main effects. δ_{ij}
- Simple calculations to estimate each component:

$$\gamma = m_{++}$$
 $\alpha_i = \frac{m_{i+}}{m_{++}}$
 $\beta_j = \frac{m_{+j}}{m_{++}}$
 $\delta_{ij} = \frac{m_{ij}}{\gamma \alpha_i \beta_j}$

• The interaction, δ_{ij} , is the ratio of observed flow to an expected flow (for the case of no interaction).

• The dis-aggregation of the components is multiplicative:

$$m_{ij} = \gamma \alpha_i \beta_j \delta_{ij}$$

- Equivalent to a saturated Poisson regression model $(R^2 = 1)$ where
 - ullet γ is constant term
 - α_i is categorical term for the origin regions
 - β_i is categorical term for the destination regions
 - δ_{ij} is an interaction term between the α_i and β_i

$$\log m_{ij} = \gamma + \alpha_i ORIG_i + \beta_j DEST_j + \delta_i ORIG_i : DEST_j$$

• When data is in a tidy format with row h would be:

$$\log y_h = \beta_0 + \beta_1 ORIG_h + \beta_2 DEST_h + \beta_3 ORIG_h : DEST_h$$

- Poisson regression models such as these where all the predictor variables are categorical - are also know as log-linear models
- Standard functions for fitting regression models, such as glm() in R will
 provide the same fitted values, but different parameter estimates
 - Use different coding system for the constraints when estimating parameters
 - Rogers' terms the parameter estimates using the equations for $\gamma, \alpha_i, \beta_j$ and δ_{ij} above the *total reference* coding system

- The migest package contains a multi_comp() function to generate parameter estimates from an origin-destination flow matrix
 - Demonstrate with previous dummy data set

```
> r <- LETTERS[1:4]
 m0 \leftarrow matrix(data = c(0, 100, 30, 70,
                        50. 0. 45. 5.
                        60, 35, 0, 40,
                        20, 25, 20, 0),
               nrow = 4, ncol = 4, byrow = TRUE,
               dimnames = list(orig = r, dest = r))
 addmargins(m0)
     dest.
orig
            B C D Sum
        0 100 30 70 200
  В
       50 0 45 5 100
  C
       60 35 0 40 135
  D
       20
           25 20
                   0 65
  Sum 130 160 95 115 500
```

```
> library(tidyverse)
 library(migest)
 mO %>%
    multi_comp() %>%
    round(3)
     dest
            Α
                                            Sum
orig
  Α
        0.000
                1.563
                         0.789
                                 1.522
                                          0.400
  В
        1.923
                0.000
                         2.368
                                 0.217
                                          0.200
  C
        1.709
                0.810
                         0.000
                                 1.288
                                          0.270
  D
        1.183
                 1.202
                         1.619
                                 0.000
                                          0.130
  Sum
        0.260
                0.320
                         0.190
                                 0.230 500.000
```

 As the model is saturated, the fitted values are the same as the observed values.

```
> multi_comp(m = m0)
     dest
                                        C
orig
                Α
                            В
                                                               Sum
  Α
        0.0000000
                    1.5625000
                                0.7894737
                                            1.5217391
                                                        0.4000000
  В
        1.9230769
                    0.0000000
                                2.3684211
                                            0.2173913
                                                        0.2000000
  C
        1.7094017
                    0.8101852
                                0.0000000
                                            1.2882448
                                                        0.2700000
  D
        1.1834320
                    1,2019231
                                1.6194332
                                            0.0000000
                                                        0.1300000
  Sum
        0.2600000
                    0.3200000
                                0.1900000
                                            0.2300000 500.0000000
>
 # fitted value for A to B
> 500 * 0.4 * 0.32 * 1.5625
[1] 100
```

- The total reference coding scheme for the parameter estimates are easier to examine than parameter estimates from a Poisson model fitted using glm() More detail on glm() in next section
- > d0 <- as.data.frame.table(x = m0, responseName = "flow")</pre> > f0 <- glm(formula = flow ~ orig + dest + orig * dest, family = poisson(),</pre> data = d0)+ > f0Call: glm(formula = flow ~ orig + dest + orig * dest, family = poisson(), data = d0)Coefficients: (Intercept) origB origC origD dest.B dest.C -24.3028.21 28.40 27.30 28.91 27.70 destD origB:destB origC:destB origD:destB origB:destC origC:destC 28.55 -57.12 -29.45-28.68-27.81 -56.10origD:destC origB:destD origC:destD origD:destD -27.70-30.85-28.96-55.85Degrees of Freedom: 15 Total (i.e. Null); O Residual Null Deviance: 463.7 Residual Deviance: 2.232e-10 AIC: 96.27

```
> # fitted and observed values are the same
> d0 %>%
    as tibble() %>%
    mutate(fit = round(f0$fitted.values, digits = 5))
# A tibble: 16 x 4
   orig dest flow
                         fit
   <fct> <fct> <dbl> <dbl>
 1 A
          Α
                    0
                           0
 2 B
          Α
                    50
                          50
 3 C
          Α
                    60
                          60
   D
          Α
                    20
                          20
 5 A
          В
                   100
                         100
 6 B
          В
                    0
                           0
 7 C
          В
                    35
                          35
 8 D
          В
                    25
                          25
 9 A
          C
                    30
                          30
10 B
                    45
                          45
11 C
          C
                    0
                           0
12 D
          C
                    20
                          20
13 A
          D
                    70
                          70
14 B
          D
                     5
                           5
15 C
          D
                          40
                    40
16 D
          D
                     0
                           0
```

- Rogers' and colleagues have used the multiplicative component model to estimate migration flow tables
- Expand to multiple dimensions
- Rectify bumpy age schedules
 - Replace reported age parameters (proportions) in the multiplicative component model with proportions from a more regular schedule.
 - Multiply the new age parameters with the existing total, origin, destination and interaction parameters to obtain new estimated flows.

Italian data in migest package

```
> italy_area
# A tibble: 3,500 x 5
  orig
            dest
                                   flow
                    year age_grp
  <chr> <chr> <chr> <chr> <dbl> <fct>
                                   <dbl>
 1 North-West North-West 1970 0-4
                                      0
2 North-East North-West 1970 0-4
                                   2350
 3 Center North-West 1970 0-4
                                   1687
4 South North-West 1970 0-4
                                   9697
5 Islands North-West 1970 0-4
                                   5139
6 North-West North-East 1970 0-4
                                   2448
7 North-East North-East 1970 0-4
                                      0
8 Center North-East 1970 0-4
                                   1063
9 South North-East 1970 0-4
                                   1560
10 Islands North-East 1970 0-4
                                    689
# ... with 3,490 more rows
# i Use `print(n = ...)` to see more rows
```

South

2.370

0.681

2.501

2.023

0.000

0.014

South

0.000

Multiplicative Component Model

```
> # single year, multiple age groups
  c0 <- italy area %>%
    filter(year == 2000) %>%
    multi_comp()
> round(c0, 3)
   age_grp = 0-4
            dest
```

```
orig
                   Center
```

Islands North-East North-West Center 0.000 1,401 Islands 0.970 0.000

0.859 1.181 1.916 0.000

North-East 1.053 North-West 0.877 2,490 1.409 0.531

> 0.016 0.007

> > 0.840

0.877

1.387

 $age_grp = 5-9$

orig

South

Center

South

Islands

North-East

North-West

Sum

dest.

Islands North-East North-West Center 0.000

1.589 1.166 0.000

1.932

2.714

0.507

0.779 1.393 0.000

0.844

1.283

0.887

1.184

0.017

0.762 1.707 0.936

0.000

1.151

0.909

1.513

1.179

0.000

1.102

0.018

2.243 0.007 0.562 0.010 2.085 0.006 1.963 0.010

Sum

0.010

0.012

0.010

0.014

0.025

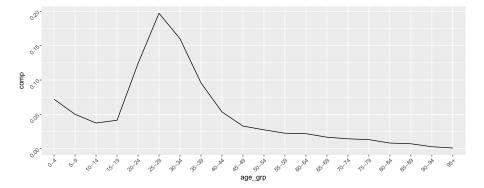
0.072

Sum

0.018

```
> # origin components (shares)
> c0 %>%
   as.data.frame.table(responseName = "comp") %>%
   filter(orig != "Sum", dest == "Sum", age_grp == "Sum")
       orig dest age_grp
                             comp
     Center Sum
                    Sum 0.1477314
    Islands Sum Sum 0.1663483
3 North-East Sum Sum 0.1245945
4 North-West Sum Sum 0.2017835
      South Sum Sum 0.3595424
5
>
 # destination components (shares)
 c0 %>%
   as.data.frame.table(responseName = "comp") %>%
   filter(orig == "Sum", dest != "Sum", age grp == "Sum")
  orig
            dest age_grp
                              comp
  Sum
      Center
                    Sum 0.23305555
  Sum
         Islands Sum 0.08368777
  Sum North-East Sum 0.25254113
  Sum North-West Sum 0.26283900
5
           South
                    Sum 0.16787656
  Sum
```

```
> # age components
> c0 %>%
+ as.data.frame.table(responseName = "comp") %>%
+ filter(orig == "Sum", dest == "Sum", age_grp != "Sum") %>%
+ ggplot(mapping = aes(x = age_grp, y = comp, group = 1)) +
+ geom_line() +
+ theme(axis.text = element_text(angle = 45, hjust = 1))
```



 Rogers' and collaborators like to shorten the multiplicative form of the log-linear model to use capital letters to represent parameters

$$m_{ij} = \gamma \alpha_i \beta_j \delta_{ij} = TO_i D_j OD_{ij}$$

 When there is multiple origin-destination tables, by different age groups, sex, education level, etc,... the notation can be easily used to study different log-linear models

$$m_{ij} = TO_i D_j A_x OD_{ij} OA_{ix}$$

• When data is in a tidy format with row h would be:

$$\log y_h = \beta_0 + \beta_1 ORIG_h + \beta_2 DEST_h + \beta_3 AGE_x + \beta_4 ORIG_h : DEST_h + \beta_5 ORIG_h : AGE_h$$

Log-Linear Models

- We can fit log-linear models in R using the glm() function (for generalised linear models)
- Requires a formula, data and family argument
- The formula argument is similar to that in xtabs(), where we use the ~ symbol to separate the the response and explanatory variables
 - For example the model in the previous slide would use formula = flow ~ orig + dest + age + orig:dest + orig:age
 - Use: or * to denote interactions
- The family argument should be set to poisson() for a log-linear model

Log-Linear Models

Example with age-specific migration flows between Italian regions in 1970

```
> d1 <- italy_area %>%
   filter(orig != dest,
          vear == 1970) %>%
   # rename so later output fits on slide
   rename(age = age_grp)
> d1
# A tibble: 400 x 5
  orig
            dest
                       year age
                                   flow
  <chr> <chr>
                       <dbl> <fct> <dbl>
 1 North-East North-West 1970 0-4
                                   2350
2 Center North-West 1970 0-4 1687
 3 South North-West 1970 0-4
                                   9697
4 Islands North-West 1970 0-4
                                   5139
5 North-West North-East 1970 0-4
                                   2448
6 Center
            North-East 1970 0-4
                                   1063
7 South North-East 1970 0-4
                                   1560
8 Islands North-East 1970 0-4
                                  689
9 North-West Center
                      1970 0-4
                                   2097
10 North-East Center 1970 0-4
                                   1183
 ... with 390 more rows
# i Use `print(n = ...)` to see more rows
```

Log-Linear Models

```
> glm(formula = flow ~ orig + dest, family = poisson(), data = d1)
Call: glm(formula = flow ~ orig + dest, family = poisson(), data = d1)
Coefficients:
  (Intercept) origIslands
                             origNorth-East origNorth-West
                                                               origSouth
      6.39791
                     0.17515
                                   -0.20852
                                                  0.99427
                                                                 0.98847
  destIslands destNorth-East destNorth-West destSouth
     -0.76940
                    -0.32536
                                   1.08367
                                                  0.02188
Degrees of Freedom: 399 Total (i.e. Null); 391 Residual
Null Deviance: 758100
Residual Deviance: 5e+05 AIC: 503100
```

- As we increase the number of dimensions of the data, it might become important to understand which dimensions of the data are most important
- We can use log-linear models with detailled migration data to
 - Understand the dominate dimensions, for example Imhoff et al. (1997) Rogers et al. (2002)
 - Predict origin-destination flows with partial data, for example Beer et al. (2010) Rogers, Willekens, and Raymer (2003) Raymer (2007)
 - Project detailed origin-destination flows, for example Raymer, Bonaguidi, and Valentini (2006)
- All the above examples involve fitting a number log-linear models based on different dimensions of the data frames
 - · Use model fit statistics to judge the best model

- One approach to choosing the most important dimensions is to fit all possible combinations of models known as *dredging* the model space
- The dredge() function in the MuMIn package will fit all combinations of regression models given an upper limit, i.e. the most complex model.
 - The number of combinations grows exponentially with the number of predictors
 - Does not allow na.action = "na.omit" set by default in glm() for handling missing values in regression models

- Fit the most complex model using glm().
 - Set na.action = na.fail to exclude failed models in when using the dredge() function later
 - Most complex model typically involves at least all two-way interactions
- The formula argument in glm() allows the use ()^2 to construct all two-way interactions, i.e. the below give the identical outputs
 - Use ()^3 for all three way interactions

1 3810+00 2 0270-02 68 123 < 20-16 ***

> f1 %>%

oriaSouth

Log-Linear Model Analysis

Models will have many estimated coefficients

Coefficients: (5 not defined because of singularities)

 Some will be non-determinable because no observations (e.g. diagonal terms such as origIslands:destIslands below) as

```
coef() %>%
   length()
Γ17 196
> summary(f1)
Call:
glm(formula = flow ~ (orig + dest + age)^2, family = poisson(),
    data = d1. na.action = na.fail)
Deviance Residuals:
     Min
                10
                      Median
                                    30
                                             Max
-12.1125 -1.4474
                      0.0186
                                1.5870
                                          8.3143
```

Estimate Std. Error z value Pr(>|z|)
(Intercept) 6.812e+00 2.085e-02 326.747 < 2e-16 ***
origIslands 4.277e-01 2.236e-02 19.126 < 2e-16 ***

origNorth-East 1.709e-01 2.390e-02 7.151 8.64e-13 ***
origNorth-West 7.906e-01 1.789e-02 44.190 < 2e-16 ***

• Pass the upper model to dredge(). Use trace = TRUE to monitor progress.

```
> library(MuMIn)
> mm <- dredge(global.model = f1, trace = TRUE)</pre>
Fixed term is "(Intercept)"
0: glm(formula = flow ~ 1, family = poisson(), data = d1, na.action = na.fail)
1 : glm(formula = flow ~ age + 1, family = poisson(), data = d1,
    na.action = na.fail)
2 : glm(formula = flow ~ dest + 1, family = poisson(), data = d1,
    na.action = na.fail)
3 : glm(formula = flow ~ age + dest + 1, family = poisson(), data = d1,
    na.action = na.fail)
4 : glm(formula = flow ~ orig + 1, family = poisson(), data = d1,
    na.action = na.fail)
5 : glm(formula = flow ~ age + orig + 1, family = poisson(), data = d1,
    na.action = na.fail)
6 : glm(formula = flow ~ dest + orig + 1, family = poisson(), data = d1,
    na.action = na.fail)
7 : glm(formula = flow ~ age + dest + orig + 1, family = poisson(),
    data = d1, na.action = na.fail)
11 : glm(formula = flow ~ age + dest + age:dest + 1, family = poisson(),
    data = d1, na.action = na.fail)
15 : glm(formula = flow ~ age + dest + orig + age:dest + 1, family = poisson(),
    data = d1. na.action = na.fail)
21 : glm(formula = flow ~ age + orig + age:orig + 1, family = poisson(),
```

delta

4271.97

8933.35

16278.99

39668.13

44748.08

49409.47

56770.72

0.00

Log-Linear Model Analysis

```
> mm
Global model call: glm(formula = flow ~ (orig + dest + age)^2, family = poisson(),
    data = d1, na.action = na.fail)
```

```
Model selection table
                                                       logLik
                                                                  AICc
```

(Int) age dst org age:dst age:org dst:org + 191

+

+

+

64 6.515 56 6.616

48 6.619 40 6.691

32 6.865 +

24 6.995 16 6.997

7.070 12 7,612

+

24 -120180.165 240411.5 233786.93

+ 115

+ 115

39

180

104

104

100

24

28

20 -160715.461 321473.1 314848.54

20 -231284.060 462610.3 455985.74

-2944.992

-5286.311

-7617,005

-11408.460

-22817.598

-25545.324

-27876.018

-31667.473

100 -114058.016 228383.6 221758.99

9 -251543.073 503104.6 496480.01 5 -306076.551 612163.3 605538.65

6624.6

10896.6

15558.0

22903.6

46292.7

51372.7

56034.1

63395.3

-82409.496 165086.5 158461.95

-86200.951 172453.1 165828.50

5 -340055.765 680121.7 673497.08 1 -380591.061 761184.1 754559.53

7.062 weight

7.684

7.325

7.734

6.398

7.012

6.653

22 7,250

39 6.019

5

- Model comparison based on model statistics measuring the goodness of fit.
 - AIC measures a goodness of fit with a penalty for the number of predictor variables.
 - AICc has a bias correction term for small samples
- Typically the origin-destination interaction term is very important for accurately predicting the age-specific origin-destination migration flows
- The time to conduct a dredging analysis increase exponentially as the number of dimensions increases.

Exercise (ex6.R)

```
# 0. a) Load the KOSTAT2022. Rproj file.
     Run the qetwd() below. It should print the directory where the
      KOSTAT2022. Rproj file is located.
getwd()
      b) Load the packages used in this exercise
library(tidyverse)
library(migest)
library(MuMIn)
##
##
##
# 1. Run the code below to read in the migration flow data for flows within the
     USA, decomposed by move type, from 6 censuses between 1940 and 2000.
us <- read_csv("./data/us_area_1940 2000.csv")
us
# 2. Show the multiplicative components, rounded to 3 digits, for the flows from
     the 2000 census
us %>%
 filter(year == 2000) %>%
  #####() %>%
 round(digits = ####)
# 3. Fit a log-linear model to the entire data set using all two-way
     interactions between the four dimensions (orig, dest, period and move type)
f <- glm(formula = flow ~ (#### + dest + #### + move type) ^####,
         family = ####(), data = us, na.action = na.fail)
```

References I

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