

Summary Migration Indices

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Background

- Compared with fertility and mortality, little attention given to the way that internal or domestic migration varies between nations.
- Comparisons of migration over time or between spatial units is complicated by many factors including:
 - Different or changing definitions of migration
 - Different or changing collection systems for migration
 - Different sizes of regions
 - Different or changing number of regions
 - Different ways of measuring distances
 - Different and changing underlying population sizes and structures
- Bell et al. (2002) brought together and proposed a number of summary measures to enable better comparisons
 - A number of these incorporated into the IMAGE software for comparisons of internal migration between many countries; Bell, Charles-Edwards, Ueffing, et al. (2015), Bell, Charles-Edwards, Kupiszewska, et al. (2015), Stillwell et al. (2016), Rees et al. (2016), Bernard, Bell, and Charles-Edwards (2014)

Background

- Bell et al. (2002) identified four main groups of migration indices:
 - Intensity of migration
 - Distance of migration
 - Migration connectivity
 - Effect of migration on the redistribution of populations

Migration intensity

- Migration intensity measures attempt to capture the overall level, or incidence, of mobility.
- Provide a single measure for comparison of migration intensities over time or space
- Some indices based on age-specific migration data
 - Will discuss later in the age-schedule section

Crude migration probability

- Crude migration intensity is a simple measure of the overall propensity to migrate
 - Similar to crude birth or death rate
- If using migration transition data, the crude migration probability (CMP) is

$$CMP = 100M/P$$

where M is the total number of migrants in a given time period and P is the population at risk

Migration intensity

- Courgeau (1973) discussed how the intensity of migration is directly related to the number of regions n in the country

$$CMP = k \log(n^2)$$

- No intrinsic meaning to a single Courgeau's k , but can be used to compare migration intensity that cannot be seen from the raw data because of differences in their zonal systems.
- Higher the value of k , greater the intensity of migration

Migration intensity

- The `index_intensity()` function in the *migest* package calculates both intensity measures, given a migration and population data
- The *migest* package also contains a data set on Korean internal migration and populations of first level administrative districts
 - Data originally downloaded from <https://kosis.kr/eng>

```
> library(tidyverse)
> library(migest)
> korea_reg
# A tibble: 2,601 x 4
  orig dest      year  flow
  <fct> <fct>    <int> <int>
1 Seoul Seoul    2012 1069300
2 Seoul Busan    2012   21437
3 Seoul Daegu    2012   13838
4 Seoul Incheon  2012   32216
5 Seoul Gwangju  2012   11811
6 Seoul Daejeon  2012   14570
7 Seoul Ulsan    2012    6799
8 Seoul Sejong   2012    1015
9 Seoul Gyeonggi-do 2012  254175
10 Seoul Gangwon-do 2012   21324
# ... with 2,591 more rows
```

Migration intensity

- The korea_pop contains the resident population in each region and year

```
> korea_pop
# A tibble: 153 x 3
  region  year population
  <chr>  <int>      <dbl>
1 Seoul   2012    10195318
2 Seoul   2013    10143645
3 Seoul   2014    10103233
4 Seoul   2015    10022181
5 Seoul   2016     9930616
6 Seoul   2017     9857426
7 Seoul   2018     9765623
8 Seoul   2019     9729107
9 Seoul   2020     9668465
10 Busan   2012     3538484
# ... with 143 more rows
```


Migration intensity

- Calculate migration and population totals in 2020

```
> m <- korea_reg %>%
+   filter(year == 2020,
+         orig != dest) %>%
+   pull(flow) %>%
+   sum()
> m
[1] 2534114
> p <- korea_pop %>%
+   filter(year == 2020) %>%
+   pull(population) %>%
+   sum()
> p
[1] 51829023
>
> index_intensity(mig_total = m, pop_total = p,
+               n = n_distinct(korea_pop$region))
# A tibble: 2 x 2
  measure      value
  <chr>        <dbl>
1 cmp          4.89
2 courceau_k   0.863
```

Migration intensity

```
> mm <- korea_reg %>%
+   group_by(year) %>%
+   filter(orig != dest) %>%
+   summarise(m = sum(flow))
> mm
# A tibble: 9 x 2
  year      m
  <int>  <int>
1  2012 2512740
2  2013 2423429
3  2014 2507796
4  2015 2551424
5  2016 2453342
6  2017 2410930
7  2018 2429184
8  2019 2384948
9  2020 2534114
```

Migration intensity

```
> pp <- korea_pop %>%
+   group_by(year) %>%
+   summarise(p = sum(population))
> pp
# A tibble: 9 x 2
  year      p
  <int>   <dbl>
1  2012 50948272
2  2013 51141463
3  2014 51327916
4  2015 51529338
5  2016 51696216
6  2017 51778544
7  2018 51826059
8  2019 51849861
9  2020 51829023
```

Migration intensity

- Passing the vectors of migration and population totals can lead to confusing output

```
> d <- mm %>%
+   left_join(pp)
> d
# A tibble: 9 x 3
   year      m      p
<int> <int> <dbl>
1  2012 2512740 50948272
2  2013 2423429 51141463
3  2014 2507796 51327916
4  2015 2551424 51529338
5  2016 2453342 51696216
6  2017 2410930 51778544
7  2018 2429184 51826059
8  2019 2384948 51849861
9  2020 2534114 51829023
```

Migration intensity

```
> index_intensity(mig_total = d$m, pop_total = d$p,
+               n = n_distinct(korea_pop$region))
# A tibble: 18 x 2
  measure      value
  <chr>      <dbl>
1 cmp        4.93
2 courceau_k 0.870
3 cmp        4.74
4 courceau_k 0.836
5 cmp        4.89
6 courceau_k 0.862
7 cmp        4.95
8 courceau_k 0.874
9 cmp        4.75
10 courceau_k 0.838
11 cmp        4.66
12 courceau_k 0.822
13 cmp        4.69
14 courceau_k 0.827
15 cmp        4.60
16 courceau_k 0.812
17 cmp        4.89
18 courceau_k 0.863
```

Migration intensity

- Set long = FALSE to put each indicator in their own column

```
> index_intensity(mig_total = d$m, pop_total = d$p,
+               n = n_distinct(korea_pop$region), long = FALSE)
# A tibble: 9 x 2
  cmp courgeau_k
  <dbl>      <dbl>
1  4.93      0.870
2  4.74      0.836
3  4.89      0.862
4  4.95      0.874
5  4.75      0.838
6  4.66      0.822
7  4.69      0.827
8  4.60      0.812
9  4.89      0.863
```

Migration intensity

- Use the `map2()` function in *purrr* to apply the function to subsets of the data
 - Provides results alongside the year
- Code below produces a nested list `i` containing the intensity measures for each year

```
> d <- mm %>%
+   left_join(pp) %>%
+   mutate(i = map2(.x = m, .y = p,
+                     .f = ~index_intensity(mig_total = .x,
+                                           pop_total = .y,
+                                           n = n_distinct(korea_pop$region),
+                                           long = FALSE)))
> d
# A tibble: 9 x 4
   year      m      p i
<int> <int> <dbl> <list>
1  2012 2512740 50948272 <tibble [1 x 2]>
2  2013 2423429 51141463 <tibble [1 x 2]>
3  2014 2507796 51327916 <tibble [1 x 2]>
4  2015 2551424 51529338 <tibble [1 x 2]>
5  2016 2453342 51696216 <tibble [1 x 2]>
6  2017 2410930 51778544 <tibble [1 x 2]>
7  2018 2429184 51826059 <tibble [1 x 2]>
8  2019 2384948 51849861 <tibble [1 x 2]>
```

Migration intensity

- The `unnest()` function in the *tidyr* package binds each component of column `i` on top of each other
 - Easier to see changes over time

```
> unnest(d, cols = i)
# A tibble: 9 x 5
```

	year	m	p	cmp	courgeau_k
	<int>	<int>	<dbl>	<dbl>	<dbl>
1	2012	2512740	50948272	4.93	0.870
2	2013	2423429	51141463	4.74	0.836
3	2014	2507796	51327916	4.89	0.862
4	2015	2551424	51529338	4.95	0.874
5	2016	2453342	51696216	4.75	0.838
6	2017	2410930	51778544	4.66	0.822
7	2018	2429184	51826059	4.69	0.827
8	2019	2384948	51849861	4.60	0.812
9	2020	2534114	51829023	4.89	0.863

Migration distance

- As migration is a spatial activity, based on movements between two locations, comparisons should take account of the way that intensities of movement vary across space.
- There are a number of measures that summarize the effects of distance across a migration system
- The distance measure between each region is not straightforward
 - Ideally measure the typical distance that migrants travel.
 - Straight line distance between population-weighted centroids of each region provide a good approximation
- The costs faced by a migrant may not be represented well by the inter-centroid distance.
 - Locations in different regions might be very close to a border, so centroids will exaggerate the distance
 - Areas can take many shapes and sizes
 - Doughnut shaped regions might have centroids not in region
 - Indented coastlines might make regions look closer than they might be (culturally, travel cost)
 - Distance measures for within region moves cause another set of problems

Average migration distance

- Summary of the average migration distance can be calculated by taking a weighted average of the migration counts, where the corresponding distances are the weights
- Bell et al. (2002) note a median average as clearly preferable to a mean average as the distribution of distances is negatively skewed, reflecting the strong distance decay effect which consistently occurs
- Comparison with the mean average distance provides a guide to the skewness

Distance decay

- A more complete method to account for the skewness in migration distances is to fit a model to predict migration counts using the distances between each regions and extract the distance parameter
- Different models could be potentially used, but tend to be based on a $\log(\text{distance})$ terms with categorical control variables for the origin and destination
- The distance decay parameter (β_3) in a Poisson log-linear model;

$$\log(m_{ij}) = \beta_0 + \beta_{1i}\text{origin} + \beta_{2j}\text{destination} + \beta_3 \log(\text{distance})$$

- The distance decay parameter of interest (β_3 in the equation above) is almost always negative, indicating an increase in migration leads to fewer predicted migrations.
 - The set of β_{1i} and β_{2j} represent some form of push and pull factors for each region (i and j)

Migration distance

- The korea_dist matrix provides estimates of the 2020 population weighted distances in kilometers between the 17 first level administrative districts in Korea
 - Data based on WorldPop estimates of Edwards et al. (2021)

```
> korea_dist
```

orig	dest				
	Busan	Chungcheongbuk-do	Chungcheongnam-do	Daegu	Daejeon
Busan	0	216	255	88	201
Chungcheongbuk-do	216	0	58	130	45
Chungcheongnam-do	255	58	0	174	54
Daegu	88	130	174	0	122
Daejeon	201	45	54	122	0
Gangwon-do	286	122	159	203	166
Gwangju	202	184	170	175	139
Gyeonggi-do	312	96	83	225	125
Gyeongsangbuk-do	107	125	176	32	128
Gyeongsangnam-do	43	190	222	72	168
Incheon	328	111	85	242	134
Jeju	306	381	368	333	337
Jeollabuk-do	201	111	96	143	66
Jeollanam-do	190	205	197	177	162
Sejong	222	37	34	141	22
Seoul	324	108	95	236	138
Ulsan	17	222	252	72	122

Migration distance

- The `index_distance()` function in the *migest* package provides three summary distance measures given a set of migration and distance measures between each origin and destination.
- The origin-destination migration flows can be given as a matrix to `m` or as a data frame, where the column names are assumed to be `orig`, `dest` and `flow`.
 - Can change using `orig_col`, `dest_col` and `flow_col` arguments
- The distance values can also be given as a matrix to `d` or as a data frame, where the column names are assumed to be `orig`, `dest` and `dist`.
- Origin and destination names in `m` and `dist` must match
- Removes all within migration moves from calculations

```
> # single year
> index_distance(m = filter(korea_reg, year == 2020),
+               d = korea_dist)
# A tibble: 3 x 2
  measure    value
  <chr>      <dbl>
1 mean      105.
2 median    68.7
3 decay     -0.852
```

Migration distance

```
> korea_reg %>%
+   nest(m = c(orig, dest, flow)) %>%
+   mutate(d = list(korea_dist)) %>%
+   mutate(i = map2(.x = m, .y = d,
+                   .f = ~index_distance(m = .x, d = .y, long = FALSE))) %>%
+   unnest(i)
```

A tibble: 9 x 6

	year	m	d	mean	median	decay
	<int>	<list>	<list>	<dbl>	<dbl>	<dbl>
1	2012	<tibble [289 x 3]>	<dbl [17 x 17]>	108.	76	-0.784
2	2013	<tibble [289 x 3]>	<dbl [17 x 17]>	107.	76	-0.795
3	2014	<tibble [289 x 3]>	<dbl [17 x 17]>	108.	76	-0.816
4	2015	<tibble [289 x 3]>	<dbl [17 x 17]>	108.	76	-0.828
5	2016	<tibble [289 x 3]>	<dbl [17 x 17]>	107.	76	-0.820
6	2017	<tibble [289 x 3]>	<dbl [17 x 17]>	108.	75.8	-0.839
7	2018	<tibble [289 x 3]>	<dbl [17 x 17]>	107.	74.8	-0.839
8	2019	<tibble [289 x 3]>	<dbl [17 x 17]>	107.	75.8	-0.836
9	2020	<tibble [289 x 3]>	<dbl [17 x 17]>	105.	68.7	-0.852

Calculating distances

- There are a number of functions to a calculate distance matrices in R
- Require a set of longitude and latitudes
- If population weighted centriods are not available from national statistics offices, a number of research centers provide estimates
 - POPGRID Data Collaborative <https://www.popgrid.org/>
- Example for Ghana using WorldPop 2020 population weighted centroids
 - CSV from <https://www.worldpop.org/doi/10.5258/SOTON/WP00703>

```
>
> g <- read_csv("../data/PWD_2020_sub_national_100m.csv") %>%
+   filter(ISO == "GHA")
> g
# A tibble: 10 x 25
   year ISO   ISO_No Country_N Adm_N GID_1 HASC   PWC_Lat PWC_Lon   Pop Density
  <dbl> <chr>   <dbl> <chr>   <chr> <chr> <chr>   <dbl>   <dbl>   <dbl>   <dbl>
1  2020 GHA     288 Ghana   Asha~ GHA.~ GH.AH   6.69   -1.60  6.43e6  258.
2  2020 GHA     288 Ghana   Bron~ GHA.~ GH.BA   7.50   -2.04  2.73e6   70.6
3  2020 GHA     288 Ghana   Cent~ GHA.~ GH.CP   5.47   -0.986 2.87e6  296.
4  2020 GHA     288 Ghana   East~ GHA.~ GH.EP   6.21   -0.517 3.12e6  188.
5  2020 GHA     288 Ghana   Grea~ GHA.~ GH.AA   5.64   -0.159 5.18e6 1414.
6  2020 GHA     288 Ghana   Nort~ GHA.~ GH.NP   9.48   -0.569 3.25e6   47
7  2020 GHA     288 Ghana   Uppe~ GHA.~ GH.UE  10.9   -0.680 1.11e6  129.
8  2020 GHA     288 Ghana   Uppe~ GHA.~ GH.UW  10.4   -2.44  8.02e5   42.2
9  2020 GHA     288 Ghana   Volta GHA.~ GH.TV   6.90    0.504 2.66e6  144.
```

Calculating distances

```
> g <- g %>%
+   filter(ISO == "GHA") %>%
+   select(Adm_N, PWC_Lon, PWC_Lat)
> g
# A tibble: 10 x 3
```

	Adm_N <chr>	PWC_Lon <dbl>	PWC_Lat <dbl>
1	Ashanti	-1.60	6.69
2	Brong Ahafo	-2.04	7.50
3	Central	-0.986	5.47
4	Eastern	-0.517	6.21
5	Greater Accra	-0.159	5.64
6	Northern	-0.569	9.48
7	Upper East	-0.680	10.9
8	Upper West	-2.44	10.4
9	Volta	0.504	6.90
10	Western	-2.16	5.45

Calculating distances

- The `dism()` function in the *geosphere* package provides great circle distance estimates in meters between centroids

```
> library(geosphere)
> ghana_dist <- g %>%
+   select(PWC_Lon, PWC_Lat) %>%
+   distm()
>
> ghana_dist
```

	[,1]	[,2]	[,3]	[,4]	[,5]	[,6]	[,7]
[1,]	0.0	102992.7	151088.53	130977.06	197276.08	329699.1	471774.1
[2,]	102992.7	0.0	254060.16	221397.05	293464.70	272448.3	399289.5
[3,]	151088.5	254060.2	0.00	97220.64	93573.16	446740.6	596619.3
[4,]	130977.1	221397.1	97220.64	0.00	74652.44	362233.8	513803.4
[5,]	197276.1	293464.7	93573.16	74652.44	0.00	427847.5	579597.5
[6,]	329699.1	272448.3	446740.58	362233.78	427847.51	0.0	151789.3
[7,]	471774.1	399289.5	596619.34	513803.44	579597.52	151789.3	0.0
[8,]	422969.2	325054.7	570615.97	511378.47	585433.81	229629.3	198393.6
[9,]	233944.5	289137.3	229081.18	136579.93	158124.96	308917.8	455693.6
[10,]	149904.5	227534.2	130136.28	200358.81	222698.68	479309.9	619145.3

	[,8]	[,9]	[,10]
[1,]	422969.2	233944.5	149904.5
[2,]	325054.7	289137.3	227534.2
[3,]	570616.0	229081.2	130136.3
[4,]	511378.5	136579.9	200358.8

Calculating distances

- Add the origin and destination names to the matrix row and columns using `dimnames()`
- Allows to combine with migration data
- Divide by 1000 to get to kilometers

```
> dimnames(ghana_dist) <- list(orig = g$Adm_N, dest = g$Adm_N)
> round(ghana_dist/1000)
```

	dest					
orig	Ashanti	Brong Ahafo	Central	Eastern	Greater Accra	Northern
Ashanti	0	103	151	131	197	330
Brong Ahafo	103	0	254	221	293	272
Central	151	254	0	97	94	447
Eastern	131	221	97	0	75	362
Greater Accra	197	293	94	75	0	428
Northern	330	272	447	362	428	0
Upper East	472	399	597	514	580	152
Upper West	423	325	571	511	585	230
Volta	234	289	229	137	158	309
Western	150	228	130	200	223	479

	dest			
orig	Upper East	Upper West	Volta	Western
Ashanti	472	423	234	150
Brong Ahafo	399	325	289	228
Central	597	571	229	130

Migration connectivity

- The size of migration flows in different migration corridors vary due to many reasons other than population sizes and distance.
 - Also reflect the strength of many other factors linking regions such as the strength of historical, cultural, social and economic ties, between regions.
- Bell et al. (2002) note a fragmentation in the literature on measures of connections in a migration system and the use of a range of terms including *spatial connectivity*, *spatial concentration*, *spatial inequality* and *spatial focusing*
- There are many indices on migration connectivity. See the *migration.indices* package for example.
- The `index_connectivity()` function provides 12 different measures, that can broadly be placed into 5 groups.
 - Requires a matrix or data frame of migration flows
 - When providing a data frame, function is assuming flows are in a column with name `flow`; change with `flow_col` if not.

Migration connectivity

```
> korea_reg %>%
+   filter(year == 2020) %>%
+   index_connectivity()
# A tibble: 11 x 2
   name                value
<chr>                <dbl>
1 connectivity         1
2 inequality_equal    0.541
3 inequality_sim      0.281
4 gini_total          0.709
5 gini_orig_standardized 0.0493
6 gini_dest_standardized 0.0517
7 mwg_orig            0.0370
8 mwg_dest            0.0389
9 mwg_mean            0.0379
10 cv                 17.9
11 acv                 3.43
```

Connectivity and Inequality

- The connectivity measure evaluates the proportion of the flows (excluding within region flows) that are non-zeros
 - More useful when many regions where populations are smaller
- Bell et al. (2002) inequality measures are based on a distributions of flows compared to distributions of expected flows
 - `inequality_equal` measures the distance of the observed flows to an expected distribution where all flows are equal
 - `inequality_sim` measures the distance of the observed flows to an expected distribution from a spatial interaction model equivalent to a Poisson regression mode for an independence fit

$$\widehat{\log(m_{ij})} = \beta_0 + \beta_{1i}\text{origin} + \beta_{2j}\text{destination}$$

- In both cases, a value of 0 shows the observed flows match the expected values (some form of equality) and 1 shows the maximum distance between the observed flows and the expected flows, i.e. maximum inequality.

Gini measures

- The Gini measures provide a value for the spatial focusing in a set of migration flow - i.e. how much of the migration is focused on a particular set of migration corridors
- Compares each flow with every other flow in the migration matrix.
- A `gini_total` value of zero indicates all flows are of equal size (no spatial focusing) to 1, only one single flow (maximum focusing).
- The `gini_orig_standardized` values provide a similar measure but compare every outflow from each origin with every other outflow from that origin.
 - Measures the extent to which the destination choices of out-migrants are spatially focused.
 - The `gini_dest_standardized` does the same but for the spatial focusing of origins of in-migrants.
 - The standardized values ensure a range of 0 and 1 - zero is no focusing

Migration totals Gini and Coefficient of Variation

- The migration weighted Gini indexes provide a measure of the focusing for the in-migration and out-migration totals (`mwg_orig` and `mwg_dest`)
- The `mwg_mean` is a simple average of `mwg_orig` and `mwg_dest` to provide a system wide measure of focusing for all migration totals.
 - As with the `gini_` measures from `index_connectivity()` values vary between zero (no focusing) and 1 (all migration goes through a single origin or destination)
- Rogers and Raymer (1998) proposed a coefficient of variation, provided by `cv` which compares the mean of the flows to the standard deviation of the flows.
 - Is not limited to 0 and 1
- The `acv` provides a similar measures of variation but based on the aggregate of coefficient of variations of in- and out-migration totals (based on the means and standard deviations of in- and out-migration totals)
 - Again, not limited to 0 and 1, but useful for comparisons across time or countries, where rising `cv` or `acv` would indicate greater inequality in migration flows or flow totals

Migration impact

- The impact of migration measures the extent to which migration acts to transform the pattern of human settlement
- Migration is already or becoming predominant mechanism leading to the redistribution of population in many regions of the world
- Descriptive studies tend to focus on regional net migration patterns
- Additional measures exist that summarize the overall effect of migration in redistributing a population across the entire system of regions

Migration effectiveness index

- The migration effectiveness index (MEI) of Shryock and Siegel (1976) compares the sum of net migration as a proportion of migration turnover, measuring the amount asymmetry or equilibrium in the migration network

$$MEI = 100 \frac{\sum_i |\text{net}|}{\sum_i \text{turnover}_i} = 100 \frac{\sum_i |m_{+i} - m_{i+}|}{\sum_i m_{+i} + m_{i+}}$$

- MEI range between 0 and 100.
- High values indicate migration is an efficient mechanism of population redistribution, based on large net totals for the given turnover.
- Values closer to zero are generated from more balanced migration systems with less population redistribution

Aggregate net migration rate

- The aggregate net migration rate (ANMR) of Bell et al. (2002) attempts to measure the overall effect of migration on the population settlement patterns by replacing the denominator of *MEI* with the each regions population P_i
- Index measures the net shift of population between regions per 100 residents in the country
 - No upper limit

$$ANMR = 100 \frac{1}{2} \frac{\sum_i |\mathbf{net}_i|}{\sum_i P_i} = 100 \frac{1}{2} \frac{\sum_i |m_{+i} - m_{i+}|}{\sum_i P_i}$$

- Product of the CMI and MEI

Preference and velocity

- The manual by United Nations Department of Economic and Social Affairs Population Division (1983) provides two other impact measures, that seem to have fallen out of favor
- The preference index is based on an expected model of migration intestines based on population shares and the overall level of migration $M \frac{p_i}{P} \frac{p_j}{P}$, where M is the total migration flow and P is the total population based on the sum of populations in each region (p_i)
- Index compares the observed flows to an expected model where migration rates in all populations are the same
 - No upper limit

$$\text{preference} = \sum_{ij} \frac{m_{ij}}{M \frac{p_i}{P} \frac{p_j}{P}}$$

- The velocity index is based on a migration velocity measure $\frac{m_{ij}}{p_i p_j}$, multiplied by the total population and summed
- Index compares observed flows to an expected models where flows sizes are determined by population sizes alone
 - No upper limit

$$\text{velocity} = \sum_{ij} \frac{m_{ij}}{p_i p_j} P$$

Migration impact

- The `index_impact()` function in *migest* calculates all four measures given a set of migration flows and population sizes in each region
 - The `p` parameters assumes column names `region` and `pop` for region and population. Change from defaults using `reg_col` and `pop_col`

```
> index_impact(
+   m = subset(korea_reg, year == 2020),
+   p = subset(korea_pop, year == 2020),
+   pop_col = "population"
+ )
# A tibble: 4 x 2
  measure      value
  <chr>        <dbl>
1 effectiveness  7.67
2 anmr          0.375
3 preference    375.
4 velocity     18.3
```

Migration impact

- Multiple years require nesting the migration and population data bases besides each other

```
> d <- korea_reg %>%
+   nest(m = c(orig, dest, flow)) %>%
+   left_join(korea_pop) %>%
+   nest(p = c(region, population))
Joining, by = "year"
> d
# A tibble: 9 x 3
   year m                                p
<int> <list>                            <list>
1  2012 <tibble [289 x 3]> <tibble [17 x 2]>
2  2013 <tibble [289 x 3]> <tibble [17 x 2]>
3  2014 <tibble [289 x 3]> <tibble [17 x 2]>
4  2015 <tibble [289 x 3]> <tibble [17 x 2]>
5  2016 <tibble [289 x 3]> <tibble [17 x 2]>
6  2017 <tibble [289 x 3]> <tibble [17 x 2]>
7  2018 <tibble [289 x 3]> <tibble [17 x 2]>
8  2019 <tibble [289 x 3]> <tibble [17 x 2]>
9  2020 <tibble [289 x 3]> <tibble [17 x 2]>
```

Migration impact

- Apply the `index_impact()` function to each row and unnest

```
> d %>%
+   mutate(i = map2(.x = m, .y = p,
+                   .f = ~index_impact(m = .x, p = .y,
+                                     pop_col = "population",
+                                     long = FALSE))) %>%
+   unnest(i)
# A tibble: 9 x 7
```

	year	m		p	effectiveness	anmr	preference	velocity
	<int>	<list>		<list>	<dbl>	<dbl>	<dbl>	<dbl>
1	2012	<tibble [289 x 3~		<tibble [17 x ~	6.07	0.300	409.	20.2
2	2013	<tibble [289 x 3~		<tibble [17 x ~	5.72	0.271	371.	17.6
3	2014	<tibble [289 x 3~		<tibble [17 x ~	5.36	0.262	434.	21.2
4	2015	<tibble [289 x 3~		<tibble [17 x ~	7.73	0.383	459.	22.7
5	2016	<tibble [289 x 3~		<tibble [17 x ~	8.47	0.402	401.	19.0
6	2017	<tibble [289 x 3~		<tibble [17 x ~	7.99	0.372	417.	19.4
7	2018	<tibble [289 x 3~		<tibble [17 x ~	9.29	0.435	398.	18.7
8	2019	<tibble [289 x 3~		<tibble [17 x ~	6.94	0.319	391.	18.0
9	2020	<tibble [289 x 3~		<tibble [17 x ~	7.67	0.375	375.	18.3

Exercise (ex3.R)

```
# 0.  a) Load the KOSTAT2021.Rproj file.
#      Run the getwd() below. It should print the directory where the
#      KOSTAT2021.Rproj file is located.
getwd()
#      b) Load the packages used in this exercise
library(tidyverse)
library(migest)
library(geosphere)
##
##
##
# 1. Run the code below to read in the bilateral data in brazil_census_tidy.csv
#      from the 1991, 2000 and 2010 Brazilian censuses
br <- read_csv("./data/brazil_census_tidy.csv")
br
# 2. Run the code below to read in the WorldPop population data for Brazil in
#      2000 and check that the orig and dest names in the br migration data match
#      the region names in br_pop
br_pop <- read_csv("./data/PWD_2000_sub_national_100m.csv",
                    locale = readr::locale(encoding = "latin1")) %>%
  filter(ISO == "BRA") %>%
  select(Adm_N, contains("PWC"), Pop)
br_pop
# check names match
unique(br$orig) %in% br_pop$Adm_N
```

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