

Estimating Net Migration

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Net migration

- At the most basic level demographers are typically interested in the net balance of migration as a component of population change.
- Might not have an interest in the complexities involved in the different scales of migration to and from each region.
- Net migration tends to be used as it is readily available.
 - Data for in- and out-migration require specialized migration question in surveys or censuses.
 - Net migration does not require any questions on migration.
- Most censuses measure population changes accurately enough in order to develop a good estimate of net migration.
- Net migration has many weaknesses for the study of migration patterns, migration trends and population projections Rogers (1990)
 - Net migrants do not exist

Net migration estimation

- Three groups of methods to derive net migration
- First two are residual methods
 - ① Vital statistics based on population change and natural increase data
 - ② Survival methods based on population change data
 - ③ Place of birth methods based on changes in migrant stock data.
- United Nations Department of Economic and Social Affairs Population Division (1983) provides a nice discussion on the relative merits of each method

Vital statistics

- The most elementary method to estimate net migration is using the demographic accounting equation

$$M = P^{t+n} - P^t - B + D$$

- Simple to calculate.
- Careful data preparation is required.
- Commonly applied to estimate net migration by sub-groups of populations where (e.g. sex)
- Less commonly applied to estimate net migration by age

$$M(x) = P^{t+n}(x + n) - P^t(x) + D(x)$$

where parenthesis represent age groups x of size n

- Not easy to accurately estimate age-specific death counts that align to period between censuses

Vital statistics

- The *migest* package has a `net_vs()` function to help obtain net migration estimates using vital statistics.
- Demonstrate using the `alabama_1970` data set in *migest*
 - Births are given in the under 10 age groups for `pop_1960`

```
> library(tidyverse)
> library(migest)
> alabama_1970
# A tibble: 68 x 6
  age_1970 sex    race pop_1960 pop_1970 us_census_sr
  <fct>    <chr>  <chr>   <dbl>   <dbl>         <dbl>
1 0-4      female white  104556  100224         0.965
2 5-9      female white  119478  115269         0.956
3 10-14     female white  120463  121922         0.997
4 15-19     female white  114627  115128         1.01
5 20-24     female white  113551  107480         0.998
6 25-29     female white   93665   87706         0.989
7 30-34     female white   76348   77285         0.996
8 35-39     female white   74278   75115         0.994
9 40-44     female white   79572   78924         0.989
10 45-49     female white   80719   78284         0.968
# ... with 58 more rows
```

Vital statsitics

- Obtain race and sex population totals
- Need to remove those not born in the original population pop_1960.

```
> d <- alabama_1970 %>%  
+   group_by(race, sex) %>%  
+   summarise(births = sum(pop_1960[1:2]),  
+             pop_1960 = sum(pop_1960) - births,  
+             pop_1970 = sum(pop_1970)) %>%  
+   ungroup()
```

``summarise()`` has grouped output by `'race'`. You can override using the ``.groups`` argument.

```
> d  
# A tibble: 4 x 5  
  race      sex    births pop_1960 pop_1970  
  <chr>   <chr>   <dbl>   <dbl>   <dbl>  
1 non-white female 126886   515483   483882  
2 non-white male  131767   467648   426452  
3 white    female 224034  1159548  1298342  
4 white    male  236481  1124061  1235489
```

Vital statistics

- Given the vital statistics `net_vs()` estimate net migration and returns three additional columns
 - `pop_change` for the population difference
 - `natural_inc` for the difference in births and deaths
 - `net` for the net migration based on the two previous columns
- The `net_vs()` function assumes `births_col = "births"` and `deaths_col = "deaths"`.
 - Can alter from default if not the case

```
> d %>%  
+   mutate(deaths = c(51449, 58845, 86880, 123220)) %>%  
+   net_vs(pop0_col = "pop_1960", pop1_col = "pop_1970")  
# A tibble: 4 x 9  
  race      sex  births pop_1960 pop_1970 deaths pop_change natural_inc    net  
  <chr>    <chr>  <dbl>   <dbl>   <dbl>   <dbl>    <dbl>    <dbl>  <dbl>  
1 non-white fema~ 126886   515483   483882   51449    -31601    75437 -107038  
2 non-white male  131767   467648   426452   58845    -41196    72922 -114118  
3 white    fema~  224034  1159548  1298342   86880    138794   137154   1640  
4 white    male  236481  1124061  1235489  123220    111428   113261  -1833
```

Difficulties

- Strictly speaking should refer to net migration estimates as a mixture of net migration and net balance of errors from the other data sources
 - Assumes international migration is nil or negligible.
 - Bogue, Hinze, and White (1982) list six difficulties with the vital statistics methods, most of which are due to the estimate is a residual from the combination of other data sources
- 1 Requires a stable administrative geography, where regions or countries do not change or at least enumerate population, births and deaths for the same units throughout the interval.
 - 2 Adjustments will be required if there has been a big change in the method to collect census, for example switching from *de jure* to *de facto* for defining place of residence
 - 3 Adjustments will be required if the birth and death periods do not align with the census dates. Typically vital statistics are annual measures starting from 1st January where as census dates are not usually on 1st January.

Difficulties

- ④ Births need to be tabulated or adjusted to mothers place of residence and deaths need to be tabulated or adjusted to place of residence or deceased. If place of occurrence is used for either then additional potential for error is created
- ⑤ Births and deaths need to be corrected for under-registration if it is known to exist.
- ⑥ Adjustments might be required to include/exclude population groups such as military or students - depending on how each are counted in the censuses and vital statistics registrations.

Survival methods

- Survival ratios can be used to compute mortality over the period, to then determine net migration as a residual.
- Survival ratios are an estimate of what proportion of a hypothetically closed population would be present at the end of the period.
 - Survival measures the force of mortality, rather than an overall population change
- Methods can be applied to total population or age-specific populations
- Preferred for age-specific net migration estimates as does not require age-specific death counts.
- Three related approaches using:
 - Forward survival ratios
 - Reverse survival ratios
 - Average survival ratios

Forward survival ratios

- Difference between the surviving expected population and observed population at the end of the period is an estimate of net migration during the interval

$$M'(x) = P^{t+n}(x+n) - s(x)P^t(x)$$

where:

- $M'(x)$ is the net migration for between t and $t+n$ for age group x
- $P^{t+n}(x+n)$ is the observed population at the end of the period ($t+n$) for age group x
- $s(x)$ is survival rate between t and $t+1$ for age group x
- $P^t(x)$ is the observed population at the start of the period (t) for age group x

Reverse survival ratios

- An alternative method is based on the reverse of the previous method
- Estimate the number of persons that would have been x years of age at the earlier census from the number who are enumerated as $x + n$ years old in the second census by applying *reverse survival ratios*

$$M''(x) = \frac{1}{s(x)} P^{t+n}(x + n) - P^t(x)$$

Average survival ratios

- The average survival ratios averages the net migration estimates from the forward and reverse survival ratios

$$\bar{M}(x) = \frac{1}{2}(M'(x) + M''(x))$$

- Siegel and Hamilton (1952) found the average survival ratio method provides the most exact approximation under normal circumstances
- Summary of assumptions for deaths:
 - Forward method: all deaths of migrants are not counted as migrants, equivalent to assuming that they all died at the place of origin.
 - Reverse method: the opposite is assumed. All migrants that die are counted as migrants, as are as those that would have moved had they survived the interval.
 - Average method: only those that died after moving are counted as migrants (approximately).

Survival methods in R

- The *migest* package contains the `net_sr()` function to calculate all three survival ratio estimates of net migration.
- Demonstrate using the `bombay_1951` data
 - Survival ratios come from a UN model life table

```
> bombay_1951
# A tibble: 13 x 5
  age_1941 age_1951 pop_1941 pop_1951 sr
  <fct>    <fct>      <dbl>   <dbl> <dbl>
1 0-4      10-14      77135   132870 0.909
2 5-9      15-19      85434   170227 0.957
3 10-14    20-24      79185   263971 0.947
4 15-19    25-29      82603   253964 0.931
5 20-24    30-34     126247   195373 0.922
6 25-29    35-39     155344   151259 0.916
7 30-34    40-44     138843   118383 0.905
8 35-39    45-49     109356    76421 0.885
9 40-44    50-54      81626    65897 0.855
10 45-49    55-59      47062    32265 0.812
11 50-54    60-64      36908    22248 0.754
12 55-59    65-69      15134     9655 0.673
13 60+      70+       25094    10100 0.387
```

Survival methods in R

```
> net_sr(bombay_1951, pop0_col = "pop_1941", pop1_col = "pop_1951")
```

```
# A tibble: 13 x 10
```

	age_1941	age_1951	pop_1941	pop_1951	sr	net_forward	net_reverse	net_average
	<fct>	<fct>	<dbl>	<dbl>	<dbl>	<dbl>	<dbl>	<dbl>
1	0-4	10-14	77135	132870	0.909	62777.	69085.	65931.
2	5-9	15-19	85434	170227	0.957	88441.	92386.	90413.
3	10-14	20-24	79185	263971	0.947	188975.	199530.	194252.
4	15-19	25-29	82603	253964	0.931	177077.	190242.	183659.
5	20-24	30-34	126247	195373	0.922	78935.	85585.	82260.
6	25-29	35-39	155344	151259	0.916	8948.	9768.	9358.
7	30-34	40-44	138843	118383	0.905	-7228.	-7990.	-7609.
8	35-39	45-49	109356	76421	0.885	-20359.	-23005.	-21682.
9	40-44	50-54	81626	65897	0.855	-3877.	-4535.	-4206.
10	45-49	55-59	47062	32265	0.812	-5959.	-7337.	-6648.
11	50-54	60-64	36908	22248	0.754	-5562.	-7382.	-6472.
12	55-59	65-69	15134	9655	0.673	-524.	-779.	-652.
13	60+	70+	25094	10100	0.387	399.	1031.	715.

```
# ... with 2 more variables: pop1_forward <dbl>, pop0_reverse <dbl>
```

Survival methods in R

- Second example using `manila_1970` where survivor ratios come from census life tables for all of the Philippines
- Births and survival rates of children are unknown

```
> manila_1970
# A tibble: 16 x 4
  age_1970 pop_1960 pop_1970 phl_census_sr
  <fct>      <dbl>    <dbl>    <dbl>
1 0-4        NA    85870      NA
2 5-9        NA    83054      NA
3 10-14      80275    79489    1.12
4 15-19      70875   101410    0.992
5 20-24      63250    90410    0.973
6 25-29      85618    56055    0.889
7 30-34      75793    44648    0.841
8 35-39      60037    36963    0.957
9 40-44      34813    28873    0.951
10 45-49     31927    23678    0.904
11 50-54     24297    19063    0.930
12 55-59     20207    14484    0.797
13 60-64     13714    10205    0.877
14 65-69      9366     6405    0.835
15 70-74      7921     3746    0.712
16 75+     11114     4779    0.562
```


Survival methods in R

- Estimate age-specific net migration for all ages, except children

```
> net_sr(manila_1970, pop0_col = "pop_1960", pop1_col = "pop_1970",  
+        survival_ratio_col = "phl_census_sr")
```

```
# A tibble: 16 x 9
```

	age_1970	pop_1960	pop_1970	phl_census_sr	net_forward	net_reverse	net_average
	<fct>	<dbl>	<dbl>	<dbl>	<dbl>	<dbl>	<dbl>
1	0-4	NA	85870	NA	0	0	0
2	5-9	NA	83054	NA	0	0	0
3	10-14	80275	79489	1.12	-10196.	-9126.	-9661.
4	15-19	70875	101410	0.992	31134.	31400.	31267.
5	20-24	63250	90410	0.973	28877.	29683.	29280.
6	25-29	85618	56055	0.889	-20082.	-22582.	-21332.
7	30-34	75793	44648	0.841	-19117.	-22723.	-20920.
8	35-39	60037	36963	0.957	-20497.	-21416.	-20957.
9	40-44	34813	28873	0.951	-4244.	-4462.	-4353.
10	45-49	31927	23678	0.904	-5189.	-5739.	-5464.
11	50-54	24297	19063	0.930	-3521.	-3788.	-3655.
12	55-59	20207	14484	0.797	-1613.	-2025.	-1819.
13	60-64	13714	10205	0.877	-1822.	-2078.	-1950.
14	65-69	9366	6405	0.835	-1417.	-1697.	-1557.
15	70-74	7921	3746	0.712	-1890.	-2657.	-2274.
16	75+	11114	4779	0.562	-1472.	-2617.	-2045.

```
# ... with 2 more variables: pop1_forward <dbl>, pop0_reverse <dbl>
```

Survival methods in R

- Estimate children net migration setting `net_children = TRUE`.
- Uses method of Shryock and Siegel (1976, p381)
 - Age 0-4: $1/4$ (ratio of 0-4 population to 15-44 female population) times net migration for females aged 15-44
 - Age 5-9: $3/4$ (ratio of 5-9 population to 20-49 female population) times net migration for females aged 20-49.
- Can alter weights in `maternal_exposure` argument
 - default is `c(0.25, 0.75)`

```
> net_sr(manila_1970, pop0_col = "pop_1960", pop1_col = "pop_1970",  
+         survival_ratio_col = "phl_census_sr", net_children = TRUE)
```

```
# A tibble: 16 x 9
```

	age_1970	pop_1960	pop_1970	phl_census_sr	net_forward	net_reverse	net_average
	<fct>	<dbl>	<dbl>	<dbl>	<dbl>	<dbl>	<dbl>
1	0-4	NA	85870	NA	-235.	-605.	-420.
2	5-9	NA	83054	NA	-8935.	-10486.	-9710.
3	10-14	80275	79489	1.12	-10196.	-9126.	-9661.
4	15-19	70875	101410	0.992	31134.	31400.	31267.
5	20-24	63250	90410	0.973	28877.	29683.	29280.
6	25-29	85618	56055	0.889	-20082.	-22582.	-21332.
7	30-34	75793	44648	0.841	-19117.	-22723.	-20920.
8	35-39	60037	36963	0.957	-20497.	-21416.	-20957.
9	40-44	34813	28873	0.951	-4244.	-4462.	-4353.
10	45-49	31927	23678	0.904	-5189.	-5739.	-5464.

Survival ratios

- The success of the above methods depend on the survival ratios.
- Ratios can be obtained from
 - Life table survival ratios (LTSR) as in `bombay_1951` example
 - Census survival ratios (CSR) as in `manila_1970` example
- Life table survival ratios are derived from the L_x columns of the life table; the ratio of persons in stationary population at age group x that are alive in comparisons to a previous age group $x - n$.

$$s_n(x) = \frac{L(x+n)}{L(x)}$$

- Can also be derived from mortality rates, if known.

Life table survival ratios

- For an accurate net migration estimate, $s_n(x)$ should
 - Measure the average mortality conditions of the period
 - Reasonably applicable to the area and population for which migration estimates are required.
- Age data to derive life tables may be inaccurate. Errors will impact the net migration estimates.

Census survival ratios

- Where appropriate life tables are not available or not appropriate, survival ratios can be computed from census age distributions
- A census survival ratio (CSR) is the ratio of the population aged $x + n$ at a given census to the population aged x at the census n years earlier.
- Computed for a nation as a whole assuming a “closed” population.
 - Adjust data for international migration before calculating CSR if international migration is a influential part of population change for a given area or group.

Age-specific birthplace data

- Example to derive the birthplace-age-specific survival ratios from the 1950 and 1960 census data, given in `usa_1960`

```
> usa_1960
# A tibble: 288 x 7
```

	birthplace	race	sex	age_1950	age_1960	pop_1950	pop_1960
	<fct>	<fct>	<fct>	<fct>	<fct>	<dbl>	<dbl>
1	New England	white	male	0-4	10-14	465097	467291
2	New England	white	female	0-4	10-14	445100	450248
3	New England	non-white	male	0-4	10-14	8419	8927
4	New England	non-white	female	0-4	10-14	8205	8896
5	New England	white	male	5-9	15-19	378265	368524
6	New England	white	female	5-9	15-19	361845	359141
7	New England	non-white	male	5-9	15-19	5421	5475
8	New England	non-white	female	5-9	15-19	5501	5977
9	New England	white	male	10-19	20-29	606335	567349
10	New England	white	female	10-19	20-29	591111	582993

```
# ... with 278 more rows
```

Age-specific birthplace data

- Focus on white males for example later on

```
> s <- usa_1960 %>%  
+   filter(sex == "male",  
+         race == "white") %>%  
+   mutate(sr = pop_1960/pop_1950) %>%  
+   select(-contains("pop"))
```

```
> s
```

```
# A tibble: 72 x 6
```

	birthplace	race	sex	age_1950	age_1960	sr
	<fct>	<fct>	<fct>	<fct>	<fct>	<dbl>
1	New England	white	male	0-4	10-14	1.00
2	New England	white	male	5-9	15-19	0.974
3	New England	white	male	10-19	20-29	0.936
4	New England	white	male	20-29	30-39	1.00
5	New England	white	male	30-39	40-49	0.996
6	New England	white	male	40-49	50-59	0.946
7	New England	white	male	50-59	60-69	0.825
8	New England	white	male	60+	70+	0.488
9	Middle Atlantic	white	male	0-4	10-14	1.01
10	Middle Atlantic	white	male	5-9	15-19	0.975

```
# ... with 62 more rows
```

Census survival ratios

- The CSR method tends to correct for systematic errors in the age data.
 - For example, get $s_n(x)$ in adolescent years greater than one as larger under-registration in 0-4 compared to 5-9 or 10-14 age groups.
- Systematic errors in the censuses might lead to survivor ratios to incorporate net census errors, that might lead to better estimate of net migration compared to LTSR.
- CSR tend to be less smooth than LTSR,
 - Perhaps more realistic age-patterns of net migration.

Limitations of census survival ratios

A number of weaknesses for CSR

- Assumes a closed population, so data must be adjusted for international migration before calculating CSR.
 - Good data on international migration data not always available
- Mortality may vary greatly in each region, so using a CSR based on national level data not always appropriate.
 - Build in regional correction factors
- Census enumeration may vary greatly in each region.
 - Build in regional correction factors
- A single census can not provide CSR for children.
 - Use birth statistics to approximate new born population for CSR calculation
 - If birth statistics are not reliable, use an approximation method using the ratio of women to children and female estimated net migration

Birthplace

- If data on lifetime migration at the start and end of the period are available, net migration can be estimated for each migrant group.
- Different procedure can be applied, depending on the availability of data
 - ① Lifetime migration totals without age characteristics
 - ② Lifetime migration data with age characteristics
- Both rely on a survival approach
 - Survival ratios are calculated by birthplace (and possibly other factors)
- If you view birthplace as just another dimension (such as sex) then the method is near identical to the survival ratio methods.
 - Can use the `net_sr()` function in `migest` once data is in correct format

Birthplace totals

- To demonstrate arranging birthplace totals with no age dimension and the application of `net_sv()` we use the `indian_sub` data in the *migest* package.

```
> indian_sub
# A tibble: 164 x 7
```

	zone	state	sex	year	in_migrants	out_migrants	net_migrants
	<chr>	<chr>	<chr>	<int>	<dbl>	<dbl>	<dbl>
1	United Prov~	United Provin~	male	1901	259836	878864	-619028
2	East Zone	East Zone	male	1901	883052	529216	353836
3	East Zone	Bihar-Orissa ~	male	1901	466126	498082	-31956
4	East Zone	Assam	male	1901	416926	31134	385792
5	Burma	Burma	male	1901	352924	4489	348435
6	South Zone	South Zone	male	1901	347416	509163	-161747
7	South Zone	Madras	male	1901	115290	450068	-334778
8	South Zone	Travancore-Co~	male	1901	42927	8515	34412
9	South Zone	Mysore	male	1901	189199	50580	138619
10	Bombay	Bombay	male	1901	311720	248149	63571

```
# ... with 154 more rows
```

Birthplace totals

- Separate columns for populations depending on birthplace
 - In state of birth or out of the state of birth.
- Rearrange data using `pivot_longer()` and `pivot_wider()` in the *tidyr* package
 - Location in its own column
 - Populations in each year in own columns
 - Work with male populations between 1921 and 1931 for those born in four selected states
 - Drop `net_migrants`, `sex` and `zone` columns

```
> d <- indian_sub %>%  
+   filter(between(year, 1921, 1931),  
+         sex == "male",  
+         state %in% c("Assam", "Madras", "Mysore", "Bombay")) %>%  
+   select(-net_migrants, -zone, -sex) %>%  
+   pivot_longer(cols = contains("migrants"), names_to = "location",  
+               values_to = "pop") %>%  
+   rename(birthplace = state)
```

Birthplace totals

```
> d
# A tibble: 16 x 4
  birthplace year location      pop
  <chr>      <int> <chr>      <dbl>
1 Assam      1921 in_migrants 671195
2 Assam      1921 out_migrants 44136
3 Madras     1921 in_migrants 97105
4 Madras     1921 out_migrants 580136
5 Mysore     1921 in_migrants 187000
6 Mysore     1921 out_migrants 45349
7 Bombay     1921 in_migrants 474553
8 Bombay     1921 out_migrants 197593
9 Assam      1931 in_migrants 754821
10 Assam     1931 out_migrants 41785
11 Madras    1931 in_migrants 119621
12 Madras    1931 out_migrants 723755
13 Mysore    1931 in_migrants 204260
14 Mysore    1931 out_migrants 54410
15 Bombay    1931 in_migrants 480557
16 Bombay    1931 out_migrants 202197
```

Birthplace totals

```
> d <- d %>%
+   mutate(location = case_when(
+     location == "in_migrants" ~ "in-state",
+     location == "out_migrants" ~ "out-of-state"
+   )) %>%
+   pivot_wider(names_from = year, values_from = pop, names_prefix = "pop_")
> d
```

A tibble: 8 x 4

	birthplace	location	pop_1921	pop_1931
	<chr>	<chr>	<dbl>	<dbl>
1	Assam	in-state	671195	754821
2	Assam	out-of-state	44136	41785
3	Madras	in-state	97105	119621
4	Madras	out-of-state	580136	723755
5	Mysore	in-state	187000	204260
6	Mysore	out-of-state	45349	54410
7	Bombay	in-state	474553	480557
8	Bombay	out-of-state	197593	202197

Birthplace totals

- Can now apply survival ratios to estimate net migration over a period
- Use a censuses survival ratio of 0.81 for both in migrants and out migrants

```
> d <- d %>%
+   mutate(sr = 0.81) %>%
+   net_sr(pop0_col = "pop_1921", pop1_col = "pop_1931")
> d
# A tibble: 8 x 10
  birthplace location      pop_1921 pop_1931      sr net_forward net_reverse
  <chr>         <chr>         <dbl>    <dbl>    <dbl>    <dbl>         <dbl>
1 Assam        in-state        671195   754821   0.81    211153.      260683.
2 Assam        out-of-state    44136    41785   0.81      6035.        7450.
3 Madras       in-state        97105   119621   0.81    40966.      50575.
4 Madras       out-of-state   580136   723755   0.81   253845.     313389.
5 Mysore       in-state       187000   204260   0.81    52790       65173.
6 Mysore       out-of-state    45349    54410   0.81    17677.      21824.
7 Bombay       in-state       474553   480557   0.81   96169.     118727.
8 Bombay       out-of-state   197593   202197   0.81   42147.      52033.
# ... with 3 more variables: net_average <dbl>, pop1_forward <dbl>,
#   pop0_reverse <dbl>
```

Birthplace totals

- To derive the net migration flow estimate for each of the states we need to make one more step
 - Subtract the net migration for the out-of-state migrants from the net migration for the in-state migrants

```
> d %>%
+   group_by(birthplace) %>%
+   summarise(net = net_forward[location == "in-state"] -
+             net_forward[location == "out-of-state"])
# A tibble: 4 x 2
  birthplace      net
  <chr>         <dbl>
1 Assam        205118.
2 Bombay        54022.
3 Madras       -212879.
4 Mysore        35113.
```


Age-specific birthplace data

- To demonstrate arranging age-specific birthplace data and the application of `net_sv()` we use the `new_england_1960` data in the *migest* package.
 - New England population totals by birthplace for white males.

```
> new_england_1960
# A tibble: 72 x 4
  birthplace      age_1960 pop_1950 pop_1960
  <fct>          <fct>      <dbl>   <dbl>
1 New England    10-14      442577  417069
2 Middle Atlantic 10-14        7651   17077
3 East North Central 10-14       1831    4376
4 West North Central 10-14        719    1313
5 South Atlantic   10-14       3451    5578
6 East South Central 10-14        679     960
7 West South Central 10-14        830    1413
8 Mountain States  10-14        533     819
9 Pacific          10-14       1730    2687
10 New England     15-19     354131  314048
# ... with 62 more rows
```

Age-specific birthplace data

- Apply the age-sex-race-birthplace specific census survivorship rate based on the US census (see previous CSR slide)

```
> d <- new_england_1960 %>%  
+   left_join(s)  
Joining, by = c("birthplace", "age_1960")  
> d  
# A tibble: 72 x 8  
  birthplace      age_1960 pop_1950 pop_1960 race  sex age_1950      sr  
  <fct>          <fct>      <dbl>   <dbl> <fct> <fct> <fct>      <dbl>  
1 New England    10-14    442577  417069 white male 0-4      1.00  
2 Middle Atlantic 10-14     7651   17077 white male 0-4      1.01  
3 East North Central 10-14     1831    4376 white male 0-4      1.01  
4 West North Central 10-14      719    1313 white male 0-4      1.00  
5 South Atlantic  10-14    3451   5578 white male 0-4      1.01  
6 East South Central 10-14      679     960 white male 0-4      1.01  
7 West South Central 10-14      830    1413 white male 0-4      1.02  
8 Mountain States  10-14      533     819 white male 0-4      1.02  
9 Pacific         10-14    1730   2687 white male 0-4      1.01  
10 New England    15-19   354131  314048 white male 5-9      0.974  
# ... with 62 more rows
```

Age-specific birthplace data

- Use the national age-sex-race-birthplace CSR to estimate net migration by birthplace and age in New England for white males

```
> d %>%
+   net_sr(pop0_col = "pop_1950", pop1_col = "pop_1960") %>%
+   relocate(contains("net"))
# A tibble: 72 x 13
```

	net_forward <dbl>	net_reverse <dbl>	net_average <dbl>	birthplace <fct>	age_1960 <fct>	pop_1950 <dbl>	pop_1960 <dbl>
1	-27596.	-27466.	-27531.	New England	10-14	442577	417069
2	9333.	9222.	9278.	Middle Atlant~	10-14	7651	17077
3	2531.	2511.	2521.	East North Ce~	10-14	1831	4376
4	594.	593.	593.	West North Ce~	10-14	719	1313
5	2086.	2062.	2074.	South Atlantic	10-14	3451	5578
6	271.	267.	269.	East South Ce~	10-14	679	960
7	567.	556.	562.	West South Ce~	10-14	830	1413
8	276.	270.	273.	Mountain Stat~	10-14	533	819
9	932.	918.	925.	Pacific	10-14	1730	2687
10	-30963.	-31782.	-31373.	New England	15-19	354131	314048

```
# ... with 62 more rows, and 6 more variables: race <fct>, sex <fct>,
#   age_1950 <fct>, sr <dbl>, pop1_forward <dbl>, pop0_reverse <dbl>
```

Exercise (ex4.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)
##
##
##
# 1. Run the code below to read in the population age structure data for Quebec
#      and a range of survival ratios
q <- read_csv("./data/quebec_1956.csv")
q
# 2. Estimate the age specific net migration counts based on the national census
#      survival ratio (column national_csr)
d1 <- #####(.data = q,
             p##### = "pop1951",
             pop1_col = ##,
             survival_ratio_col = #####)
d1
# 3. Find the total net migration estimates for the net_average method for the
#      estimates in the previous question
#####(d1$net_average)
# 4. Estimate the age specific net migration counts based on the Quebec life
```

References

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