## Bonus exercise: Replicating Nishikido, Cui and Esteve

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In Exercise 1, we used the Kitagawa method to decompose the general fertility rate by age. However, this method can be used to decompose any crude rate along any dimension. In this exercise we will use the Kitagawa decomposition to analyse differences in TFR. Moreover, we will decompose along two dimensions: age and partnership status. This exercise is based on Nishikido, Cui and Esteve (2022). The data have been simulated for privacy reasons.

The elements we need are the same as in Exercise 1:

- $t_1$  is the initial period and  $t_2$  is the final period
- $b_{ix}$  = number of births for age i and group x
- $ASFR_x$  = age-specific fertility rate for group x
- $N_{ix}$  = is the mid-year population for age i and group x
- N =total population

We know that  $TFR_x = \sum_x ASFR(x)$  and that the difference between two TFRs can be expressed as

$$\Delta TFR = \sum_{x} \sum_{p} ASFR(x, p, t = t_2) \times \frac{N(x, p, t = t_2)}{N(p, t = t_2)} - \sum_{x} \sum_{p} ASFR(x, p, t = t_1) \times \frac{N(x, p, t = t_1)}{N(p, t = t_1)} \quad (1)$$

Note the double summation, one for age and one for partnership status. Because we are decomposing along two dimensions, we need a double summation also in the Kitagawa formula:

$$\Delta TFR = \sum_{x} \sum_{p} (ASFR(x, p, t = t_2) - ASFR(x, p, t = t_1)) \times (\frac{N(x, p, t = t_2)}{N(p, t = t_2)} + \frac{N(x, p, t = t_1)}{N(p, t = t_1)}) \times 0.5 + \sum_{x} \sum_{p} (ASFR(x, p, t = t_2) + ASFR(x, p, t = t_1)) \times 0.5 \times (\frac{N(x, p, t = t_2)}{N(p, t = t_2)} - \frac{N(x, p, t = t_1)}{N(p, t = t_1)})$$

NB: In this case GFR and TFR are identical, because we consider a birth cohort with no observed mortality. Thus, total exposures do not change by age.

The file "NishikidoCuiEsteve\_Kitagawa.RData" contains simulated data on births and exposures for women born between 1965 and 1969 in Spain and Sweden by age and partnership status (kindly supplied by the authors). The exercise consists in decomposing the changes in TFR following Kitagawa. Once again, the aim is to disentangle the contribution of differences in rates and the contribution of differences in the composition of the population.

Start by loading the data and the tidyverse package.

load('NishikidoCuiEsteve\_Kitagawa.RData')

library(tidyverse)

Then select births and exposures for each of the countries and store them in vectors.

```
# Select births for Spain, for unpartnered and partnered women
bx0 ESP <- data %>%
  filter(pop=="ESP", partnered==0) %>%
  pull(births)
bx1_ESP <- data %>%
  filter(pop=="ESP", partnered==1) %>%
  pull(births)
# Select population for Spain, for unpartnered and partnered women
Nx0_ESP <- data %>%
  filter(pop=="ESP", partnered==0) %>%
  pull(exposure)
Nx1_ESP <- data %>%
  filter(pop=="ESP", partnered==1) %>%
  pull(exposure)
# Do the same for Sweden
bx0_SWE <- data %>%
  filter(pop=="SWE", partnered==0) %>%
  pull(births)
bx1_SWE <- data %>%
  filter(pop=="SWE", partnered==1) %>%
  pull(births)
NxO_SWE <- data %>%
  filter(pop=="SWE", partnered==0) %>%
  pull(exposure)
Nx1_SWE <- data %>%
  filter(pop=="SWE", partnered==1) %>%
  pull(exposure)
```

First, we look at the difference in TFRs.

```
# Spain
TFR_ESP <- sum(bx1_ESP+bx0_ESP)/(Nx1_ESP+Nx0_ESP)[1]
# the same for Sweden
TFR_SWE <- sum(bx1_SWE+bx0_SWE)/(Nx1_SWE+Nx0_SWE)[1]
# difference in TFR
(diff <- TFR_SWE - TFR_ESP)</pre>
```

## ## [1] 0.1306009

Let's decompose it and see if the difference in the fertility rate between Spain and Sweden is due to a composition effect in the population or to differences in the partnership- and age-specific fertility rates.

We need to calculate the partnership- and age-specific fertility rates, which are the number of births per subpopulation divided by the exposure for both countries. We also need the exposures, by country, partnership status and age.

```
# Partnership-specific ASFR for Spain
ASFRO_ESP <- bx0_ESP/Nx0_ESP
ASFR1_ESP <- bx1_ESP/Nx1_ESP
# And Sweden
ASFRO_SWE <- bx0_SWE/Nx0_SWE
ASFR1_SWE <- bx1_SWE/Nx1_SWE</pre>
```

Now, we calculate the Kitagawa components. They need to be specific to each age and partnership status combination.

```
# Unpartnered women

RCO <- 0.5*(Nx0_SWE/(Nx0_SWE+Nx1_SWE) + Nx0_ESP/(Nx0_ESP+Nx1_ESP))*(ASFRO_SWE-ASFRO_ESP)

CCO <- 0.5*(ASFRO_SWE+ASFRO_ESP)*(Nx0_SWE/(Nx0_SWE+Nx1_SWE)-Nx0_ESP/(Nx0_ESP+Nx1_ESP))

# Partnered women

RC1 <- 0.5*(Nx1_SWE/(Nx0_SWE+Nx1_SWE) + Nx1_ESP/(Nx0_ESP+Nx1_ESP))*(ASFR1_SWE-ASFR1_ESP)

CC1 <- 0.5*(ASFR1_SWE+ASFR1_ESP)*(Nx1_SWE/(Nx0_SWE+Nx1_SWE)-Nx1_ESP/(Nx0_ESP+Nx1_ESP))
```

Then, we sum across partnership status. This is the first (most internal) summation.

```
# Sum across partnership status (first summation)
RC_age <- RC0+RC1
CC_age <- CC0+CC1</pre>
```

Then, we sum across ages, to get the total rate and composition effects.

```
# Sum across ages (second summation)
RC <- sum(RC_age)
CC <- sum(CC_age)</pre>
```

Let's check the results

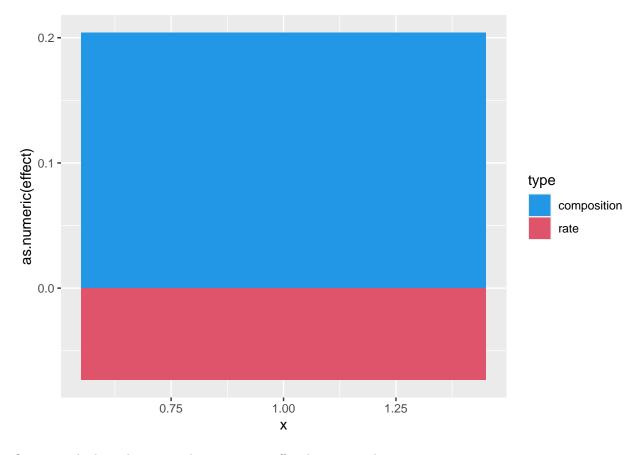
RC+CC

```
## [1] 0.1306009
diff
```

```
## [1] 0.1306009
```

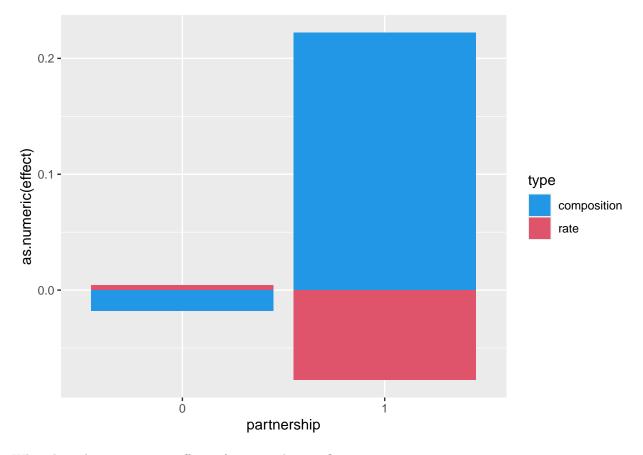
We can plot these results in many different ways. We can simply look at the total rate and composition effects.

```
cbind(type="rate",effect=sum(RCO+RC1)) %>%
    rbind(cbind(type="composition",effect=sum(CCO+CC1))) %>%
    as.data.frame() %>%
    ggplot() +
    geom_bar(aes(x=1, y=as.numeric(effect), fill=type), stat="identity", position="stack") +
    scale_fill_manual(values=c(4,2))
```



Or we can look at the rate and composition effect by partnership status.

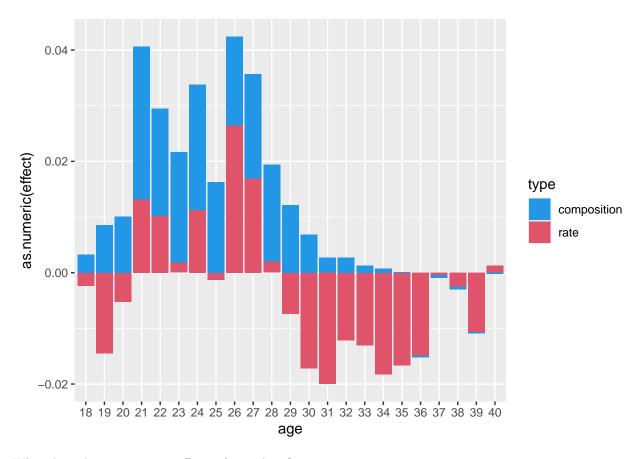
```
cbind(partnership=0,type="rate",effect=sum(RC0)) %>%
   rbind(cbind(partnership=1,type="rate",effect=sum(RC1))) %>%
   rbind(cbind(partnership=0,type="composition",effect=sum(CC0))) %>%
   rbind(cbind(partnership=1,type="composition",effect=sum(CC1))) %>%
   as.data.frame() %>%
   ggplot() +
   geom_bar(aes(x=partnership, y=as.numeric(effect), fill=type), stat="identity", position="stack") +
   scale_fill_manual(values=c(4,2))
```



What does the composition effect refer to, in this case?

Finally, we can get the contributions by age, divided in rate and composition effects.

```
cbind(age=18:40,type="rate",effect=RC_age) %%
  rbind(cbind(age=18:40,type="composition",effect=CC_age)) %>%
  as.data.frame() %>%
ggplot() +
  geom_bar(aes(x=age, y=as.numeric(effect), fill=type), stat="identity", position="stack") +
  scale_fill_manual(values=c(4,2))
```



What does the composition effect refer to, here?