

Social Inclusion Analysis

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```
dat <- read_csv("~/urbanplanning/preliminary_data.csv")

## Rows: 169989 Columns: 21

## -- Column specification -----
## Delimiter: ","
## chr (2): ethnicity, group
## dbl (19): participant, like.current.city, natives.like.me, natives.lookdown...

##
## i Use `spec()` to retrieve the full column specification for this data.
## i Specify the column types or set `show_col_types = FALSE` to quiet this message.

dat$migration.type <- dat$migration.self
dat$migration.type[dat$migration.type == 1] <- "self"
dat$migration.type[dat$migration.fellows == 1] <- "fellows"
dat$migration.type[dat$migration.relatives == 1] <- "relatives"

col_names <- c("ethnicity", "marriage", "diabete_hypertension", "group", "recent_disease", "insuranced", "gender")
dat[, col_names] <- lapply(dat[, col_names], factor)

dat$income.normalized <- (dat$income - mean(dat$income, na.rm = TRUE)) / sd(dat$income, na.rm = TRUE)
dat$income.thousand <- dat$income / 1000

dat$participant <- as.character(dat$participant)

dat$health <- as.numeric(dat$health)

head(dat)

## # A tibble: 6 x 24
##   participant like.current.city natives.like.me natives.lookdown~ native.customs.~
##   <chr>          <dbl>          <dbl>          <dbl>          <dbl>
## 1 0              1             -1              1             -1
## 2 1              2              1              1             -1
## 3 2              2              1              1              1
## 4 3              1              1             -1             -1
## 5 4              1              1             -2              1
## 6 5              2              1             -2              2
## # ... with 19 more variables: i.am.native <dbl>, income <dbl>, education <dbl>,
## #   gender <fct>, ethnicity <fct>, marriage <fct>, worked_before5.1 <fct>,
## #   diabete_hypertension <fct>, recent_disease <fct>, group <fct>,
## #   participated_in_group_activity <dbl>, insuranced <fct>,
## #   migration.self <dbl>, migration.relatives <dbl>, migration.fellows <dbl>,
## #   health <dbl>, migration.type <chr>, income.normalized <dbl>,
```

```

## #   income.thousand <dbl>

dat$inclusion <- dat$like.current.city + dat$native.like.me - dat$native.lookdown.me - dat$native.customs.better
dat$inclusion <- (dat$inclusion - min(dat$inclusion, na.rm = TRUE)) / (max(dat$inclusion, na.rm = TRUE) - min(dat$inclusion, na.rm = TRUE))
dat$loneliness.level <- -dat$inclusion + dat$migration.self - dat$migration.relative - dat$migration.fellows
dat$loneliness.level <- (dat$loneliness.level - min(dat$loneliness.level, na.rm = TRUE)) / (max(dat$loneliness.level, na.rm = TRUE) - min(dat$loneliness.level, na.rm = TRUE))

dat

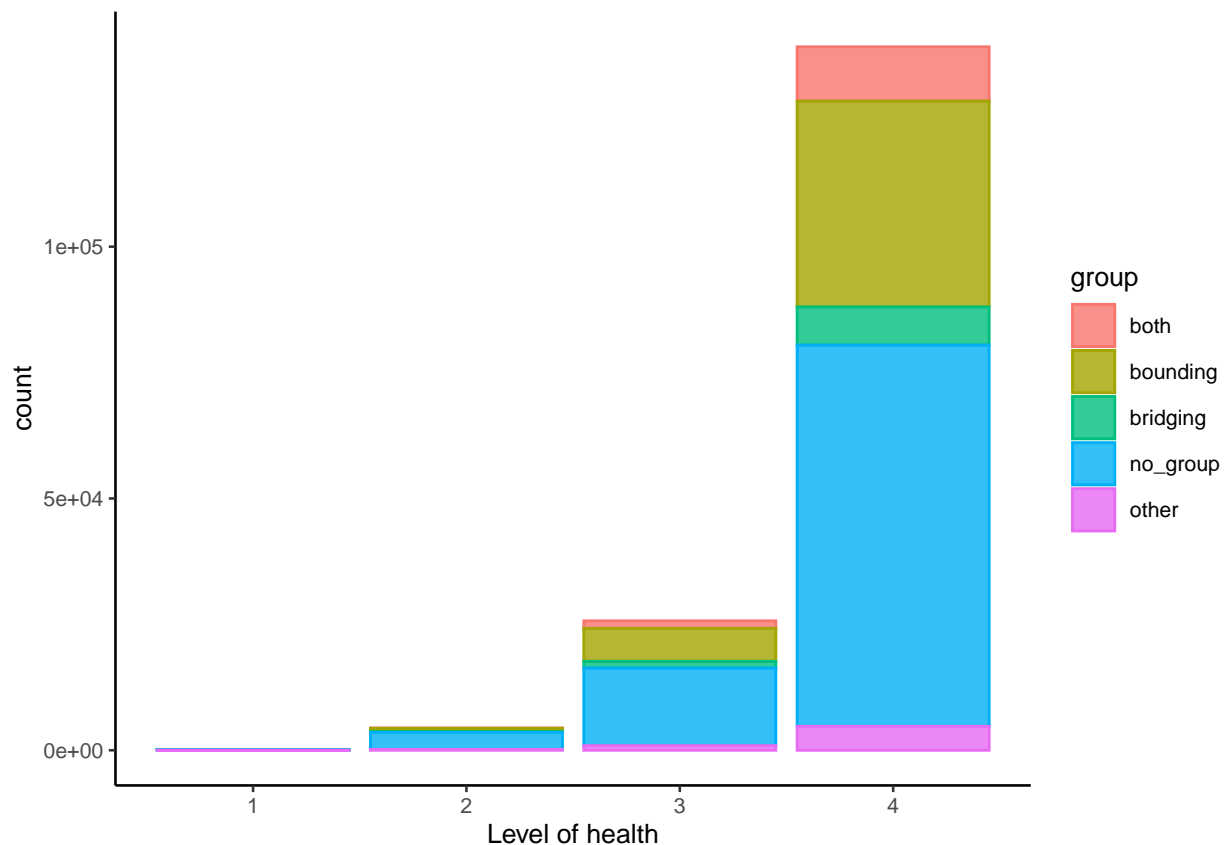
## # A tibble: 169,989 x 26
##   participant like.current.city native.like.me native.lookdown.me
##   <chr>           <dbl>           <dbl>           <dbl>
## 1 0               1             -1              1
## 2 1               2              1              1
## 3 2               2              1              1
## 4 3               1              1             -1
## 5 4               1              1             -2
## 6 5               2              1             -2
## 7 6               1              1             -2
## 8 7               1              1             -2
## 9 8               2              1             -1
## 10 9              2              2             -2
## # ... with 169,979 more rows, and 22 more variables:
## #   native.customs.better <dbl>, i.am.native <dbl>, income <dbl>,
## #   education <dbl>, gender <fct>, ethnicity <fct>, marriage <fct>,
## #   worked_before5.1 <fct>, diabete_hypertension <fct>, recent_disease <fct>,
## #   group <fct>, participated_in_group_activity <dbl>, insured <fct>,
## #   migration.self <dbl>, migration.relative <dbl>, migration.fellows <dbl>,
## #   health <dbl>, migration.type <chr>, income.normalized <dbl>, ...

cbPalette <- c("#e61212", "#ffb300", "#22ff00", "#0015ff", "#00fbff")

p1 <- ggplot(dat, aes(health, colour = group, fill = group)) +
  geom_bar(alpha = 0.8) +
  labs(
    x = "Level of health",
    colour = "group") +
  theme_classic(base_size = 10)

p1

```



```
m0 <- lmer(health ~ group + like.current.city + natives.like.me + natives.lookdown.me + native.customs.l
+ migration.type + (1|income) + (1|worked_before5.1) + (1| education) + (1 |gender) + (1|ethnicity)
data=dat)
```

```
m1 <- lmer(health ~ group + like.current.city + natives.like.me + natives.lookdown.me + native.customs.l
+ migration.type + (1|income.normalized) + (1|worked_before5.1) + (1| education) + (1 |gender) + (1|ethnicity)
data=dat)
```

```
m2 <- lmer(health ~ group + inclusion + participated_in_group_activity + migration.type
+ (1|income.normalized) + (1|worked_before5.1) + (1| education) + (1 |gender) + (1|ethnicity)
data=dat)
```

```
m3 <- lmer(health ~ group + inclusion + participated_in_group_activity + loneliness.level
+ (1|income.normalized) + (1|worked_before5.1) + (1| education) + (1 |gender) + (1|ethnicity)
data=dat)
```

```
screenreg(c(m0,m1,m2,m3))
```

```
##
## =====
##               Model 1      Model 2      Model 3      Model 4
## -----
## (Intercept)      3.55 ***      3.55 ***      3.52 ***      3.43 ***
##                (0.14)      (0.14)      (0.14)      (0.14)
## groupbounding      0.00      0.00      0.00      0.00
##                (0.00)      (0.00)      (0.00)      (0.00)
## groupbridging    -0.03 ***    -0.03 ***    -0.03 ***    -0.03 ***
##                (0.01)      (0.01)      (0.01)      (0.01)
```

## groupno_group	-0.03 ***	-0.03 ***	-0.03 ***	-0.03 ***
##	(0.00)	(0.00)	(0.00)	(0.00)
## groupother	-0.03 ***	-0.03 ***	-0.03 ***	-0.03 ***
##	(0.01)	(0.01)	(0.01)	(0.01)
## like.current.city	0.02 ***	0.02 ***		
##	(0.00)	(0.00)		
## natives.like.me	0.00	0.00		
##	(0.00)	(0.00)		
## natives.lookdown.me	-0.01 ***	-0.01 ***		
##	(0.00)	(0.00)		
## native.customs.better	-0.00	-0.00		
##	(0.00)	(0.00)		
## i.am.native	-0.01 ***	-0.01 ***		
##	(0.00)	(0.00)		
## participated_in_group_activity	-0.01 ***	-0.01 ***	-0.01 ***	-0.01 ***
##	(0.00)	(0.00)	(0.00)	(0.00)
## migration.typefellows	-0.05 *	-0.05 *	-0.04 *	
##	(0.02)	(0.02)	(0.02)	
## migration.typerelatives	-0.07 ***	-0.07 ***	-0.06 ***	
##	(0.02)	(0.02)	(0.02)	
## migration.typeself	-0.04	-0.04	-0.03	
##	(0.02)	(0.02)	(0.02)	
## inclusion			0.09 ***	0.10 ***
##			(0.01)	(0.01)
## loneliness.level				0.06 ***
##				(0.00)
## -----				
## AIC	201862.82	201862.82	202028.27	202033.50
## BIC	202093.53	202093.53	202218.86	202204.03
## Log Likelihood	-100908.41	-100908.41	-100995.14	-100999.75
## Num. obs.	167873	167873	167873	167873
## Num. groups: income	654			
## Num. groups: ethnicity	19	19	19	19
## Num. groups: education	7	7	7	7
## Num. groups: marriage	6	6	6	6
## Num. groups: recent_disease	3	3	3	3
## Num. groups: insured	2	2	2	2
## Num. groups: gender	2	2	2	2
## Num. groups: worked_before5.1	2	2	2	2
## Var: income (Intercept)	0.06			
## Var: ethnicity (Intercept)	0.00	0.00	0.00	0.00
## Var: education (Intercept)	0.01	0.01	0.01	0.01
## Var: marriage (Intercept)	0.02	0.02	0.02	0.02
## Var: recent_disease (Intercept)	0.02	0.02	0.02	0.02
## Var: insured (Intercept)	0.00	0.00	0.00	0.00
## Var: gender (Intercept)	0.00	0.00	0.00	0.00
## Var: worked_before5.1 (Intercept)	0.01	0.01	0.01	0.01
## Var: Residual	0.19	0.19	0.19	0.19
## Num. groups: income.normalized		654	654	654
## Var: income.normalized (Intercept)		0.06	0.06	0.06
## =====				
## *** p < 0.001; ** p < 0.01; * p < 0.05				