

# Multinomial regression of spouse characteristics for the ChitwanABM

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Analyzes the relationship grid data from the CVFS to build a multinomial regression predict the probability of marrying a spouse within each of a set of age groups, together with the probability of marrying someone of a different ethnicity.

The results of this analysis are used to inform the marriage process in the ChitwanABM. Once the determination has been made that a woman is marrying (based on the probability derived from event history analysis), the multinomial regression developed here is used to assign a probability of marriage to each possible spouse, based on the age of that person, so that the age differential between spouses is realistic.

## Load the data and setup R

```
library(ggplot2)
library(mlogit)
```

```
## Loading required package: Formula
```

```
## Loading required package: statmod
```

```
## Loading required package: lmtest
```

```
## Loading required package: zoo
```

```
## Attaching package: 'zoo'
```

```
## The following object(s) are masked from 'package:base':
##
## as.Date, as.Date.numeric
```

```
## Loading required package: maxLik
```

```
## Loading required package: miscTools
```

```
## Loading required package: MASS
```

```
library(arm)
```

```
## Loading required package: Matrix
```

```
## Loading required package: lattice
```

```
## Loading required package: lme4
```

```
## Attaching package: 'lme4'
```

```
## The following object(s) are masked from 'package:stats':  
##  
## AIC, BIC
```

```
## Loading required package: R2WinBUGS
```

```
## Loading required package: coda
```

```
## Attaching package: 'coda'
```

```
## The following object(s) are masked from 'package:lme4':  
##  
## HPDinterval
```

```
## Loading required package: abind
```

```
## Loading required package: foreign
```

```
## arm (Version 1.5-05, built: 2012-6-6)
```

```
## Working directory is  
## C:/users/azvoleff/code/r/chitwan_r_files/Event_History_Analysis
```

```
## Attaching package: 'arm'
```

```
## The following object(s) are masked from 'package:coda':  
##  
## traceplot
```

```
library(rms) # Note 'Design' package was renamed to 'rms'
```

```
## Loading required package: Hmisc
```

```
## Loading required package: survival
```

```
## Loading required package: splines
```

```
## Hmisc library by Frank E Harrell Jr
##
## Type library(help='Hmisc'), ?Overview, or ?Hmisc.Overview') to see overall
## documentation.
##
## NOTE:Hmisc no longer redefines [.factor to drop unused levels when
## subsetting. To get the old behavior of Hmisc type dropUnusedLevels().
```

```
## Attaching package: 'Hmisc'
```

```
## The following object(s) are masked from 'package:survival':
##
## untangle.specials
```

```
## The following object(s) are masked from 'package:base':
##
## format.pval, round.POSIXt, trunc.POSIXt, units
```

```
## Attaching package: 'rms'
```

```
## The following object(s) are masked from 'package:survival':
##
## Surv
```

```
## The following object(s) are masked from 'package:lmtest':
##
## lrtest
```

```
theme_update(theme_bw(base_size = 10))
```

```
load("V:/Nepal/CVFS_R_format/hhrel_with_respIDs.Rdata")
hhrel <- hhrel_with_respIDs

# Drop 'other' ethnicity for consistency with existing work
hhrel <- hhrel[!(hhrel$ethnic == "Other"), ]
hhrel$ethnic <- factor(hhrel$ethnic)
hhrel$CENGENDR <- factor(hhrel$CENGENDR)

hhrel$age_cat <- cut(hhrel$CENAGE, breaks = c(0, 15, 20, 30, 40,
      50, 60, 999), ordered_result = TRUE)
```

## Basic statistics

First look at some basic statistics on who is married, and on how many spouses they have, by gender.

```
hhrel$HASSPOUSE1 <- !is.na(hhrel$SPOUSE1)
hhrel$HASSPOUSE2 <- !is.na(hhrel$SPOUSE2)
hhrel$HASSPOUSE3 <- !is.na(hhrel$SPOUSE3)
xtabs(~CENGENDR + HASSPOUSE1, data = hhrel)
```

```
##          HASSPOUSE1
## CENGENDR FALSE TRUE
##   female  2218 1892
##   male    2293 1849
```

```
xtabs(~CENGENDR + HASSPOUSE2, data = hhrel)
```

```
##          HASSPOUSE2
## CENGENDR FALSE TRUE
##   female  4110    0
##   male    4109   33
```

```
xtabs(~CENGENDR + HASSPOUSE3, data = hhrel)
```

```
##          HASSPOUSE3
## CENGENDR FALSE TRUE
##   female  4110    0
##   male    4140    2
```

Now look at who has more than one spouse (only males do) by age group:

```
xtabs(~age_cat + HASSPOUSE2, data = hhrel)
```

```
##          HASSPOUSE2
## age_cat  FALSE TRUE
## (0,15]    3228    0
## (15,20]    989    0
## (20,30]   1393    1
## (30,40]    970    7
## (40,50]    711    7
## (50,60]    470   10
## (60,999]   458    8
```

```
xtabs(~age_cat + HASSPOUSE3, data = hhrel)
```

```
##          HASSPOUSE3
## age_cat  FALSE TRUE
## (0,15]    3228    0
## (15,20]    989    0
## (20,30]   1394    0
## (30,40]    977    0
## (40,50]    718    0
## (50,60]    479    1
## (60,999]   465    1
```

Having multiple wives is mostly confined to older men. Now look into the difference in spouse age, by gender.

```
spouse1_row <- match(hhrel$SPOUSE1, hhrel$RESPID)
hhrel$sp_age <- hhrel$CENAGE[spouse1_row]
```

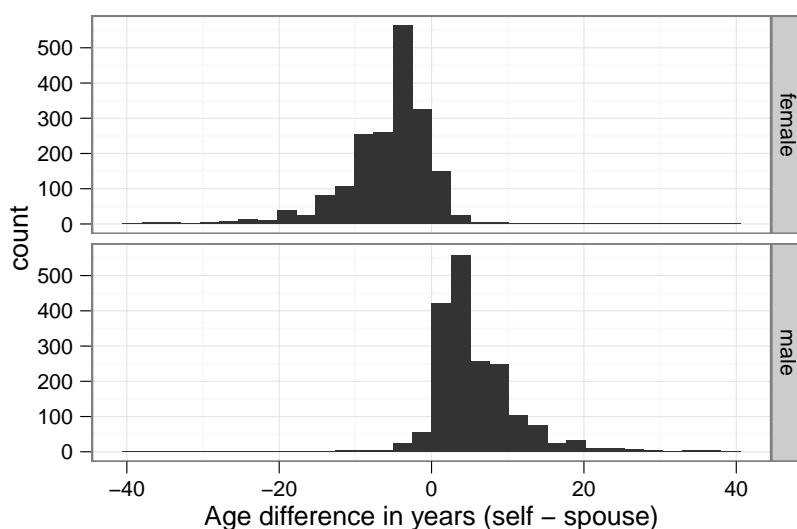
```
hhrel$sp_age_cat <- hhrel$age_cat[spouse1_row]
hhrel$sp_age_diff <- hhrel$CENAGE - hhrel$sp_age
xtabs(~age_cat + sp_age_cat, data = hhrel)
```

```
##           sp_age_cat
## age_cat  (0,15] (15,20] (20,30] (30,40] (40,50] (50,60] (60,999]
## (0,15]      0      1      2      0      0      0      0
## (15,20]     1     88    160      8      0      1      0
## (20,30]     2    160    689    271     27      6      1
## (30,40]     0      7    265    425    194     31      8
## (40,50]     0      0     26    189    268    146     41
## (50,60]     0      1      4     28    142    133     85
## (60,999]    0      0      1      7     38     83    188
```

```
qplot(sp_age_diff, facets = CENGENDR ~ ., xlab = "Age difference in years (self - spouse)",
      data = hhrel)
```

```
## stat_bin: binwidth defaulted to range/30. Use 'binwidth = x' to adjust
## this.
```

```
## stat_bin: binwidth defaulted to range/30. Use 'binwidth = x' to adjust
## this.
```



*Comparison of spouse age difference by gender*

Setup some age categories for the difference in spouse age.

```
hhrel$sp_age_diff_cat <- cut(hhrel$sp_age_diff, breaks = c(-999,
  -15, -10, -5, -2, -1, 0, 1, 2, 5, 10, 15, 999), ordered_result = TRUE)
xtabs(~sp_age_diff_cat + age_cat + CENGENDR, data = hhrel)
```

```
## , , CENGENDR = female
##
```

```
##           age_cat
## sp_age_diff_cat (0,15] (15,20] (20,30] (30,40] (40,50] (50,60] (60,999]
## (-999,-15]      0      4      29      34      41      14      2
## (-15,-10]       1     17     64     68     59     30     12
## (-10,-5]        1     79    242    146     95     51     30
## (-5,-2]         1     78    219    122     77     23     23
## (-2,-1]         0     25     48     24     14     12      7
## (-1,0]          0      4     27     30     16     11      8
## (0,1]           0      3      8      8      5      6      3
## (1,2]           0      0      5      6      5      2      3
## (2,5]           0      1      3      5      8      3      5
## (5,10]          0      0      1      0      4      1      2
## (10,15]         0      0      0      0      1      0      2
## (15,999]        0      0      0      0      1      0      0
##
## , , CENGENDR = male
##
##           age_cat
## sp_age_diff_cat (0,15] (15,20] (20,30] (30,40] (40,50] (50,60] (60,999]
## (-999,-15]      0      0      1      0      0      0      0
## (-15,-10]       0      0      1      2      0      0      3
## (-10,-5]        0      0      1      3      6      1      3
## (-5,-2]         0      2      9      9     10      4      4
## (-2,-1]         0      3      9      8      4      6      3
## (-1,0]          0      6     27     30     14     12      8
## (0,1]           0     17     54     25     13     14      7
## (1,2]           0      8     81     42     36     13     14
## (2,5]           0     11    212    162    100     45     28
## (5,10]          0      0    106    165     99     75     60
## (10,15]         0      0      9     33     49     50     37
## (15,999]        0      0      0      8     13     20     53
##
```

Do people marry outside of their ethnic group?

```
spouse1_row <- match(hhrel$SPOUSE1, hhrel$RESPID)
hhrel$spouse_ethnicity <- hhrel$ethnic[spouse1_row]
xtabs(~ethnic + spouse_ethnicity, data = hhrel)
```

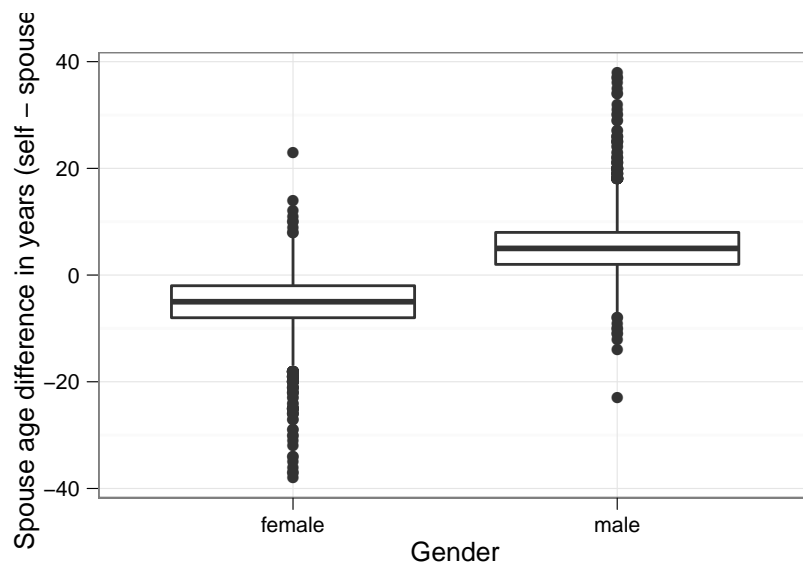
```
##           spouse_ethnicity
## ethnic      UpHindu HillTibeto LowHindu Newar TeraiTibeto
## UpHindu      1733         2         0         0         0
## HillTibeto    2         633         0         0         2
## LowHindu      0         0        413         0         0
## Newar         0         0         0        244         0
## TeraiTibeto   0         2         0         0        696
```

Marriages outside of your ethnic group are VERY uncommon. There are only 4 in the data. So we will disallow these marriages in the model - there are not enough of them to develop any kind of predictive model of when they might occur.

Make a few final summary plots: the mean of spouse\_age\_diff versus gender, and a histogram of spouse\_age\_diff by gender.

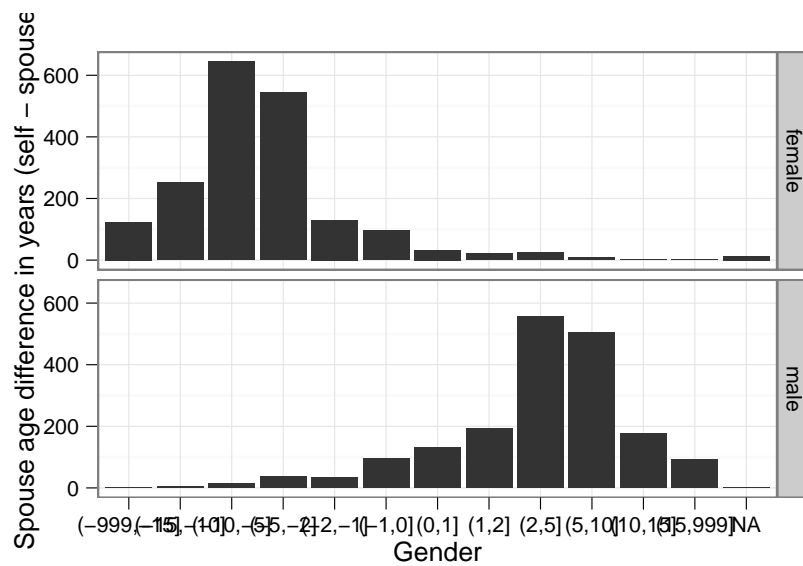
```
qplot(CENGENDR, sp_age_diff, geom = "boxplot", xlab = "Gender", ylab = "Spouse age difference in years (self - spouse)",
      data = hhrel)
```

```
## Warning: Removed 4525 rows containing non-finite values (stat_boxplot).
```



*plot of chunk mean-sp-age-diff-versus-age*

```
ggplot(sp_age_diff_cat, facets = CENGENDR ~ ., geom = "histogram",
  xlab = "Gender", ylab = "Spouse age difference in years (self - spouse)",
  data = hhrel[hhrel$HASSPOUSE1, ])
```



*Histogram of marriages by spouse age categories and gender*

# Multinomial logistic regression predicting spouse age difference based on gender

```
hhrel_mlogit_data <- mlogit.data(hhrel, varying = NULL, choice = "sp_age_diff_cat",
  shape = "wide")
mlogit_spouseage <- mlogit(sp_age_diff_cat ~ 1 | CENGENDR, data = hhrel_mlogit_data)
summary(mlogit_spouseage)
```

```
##
## Call:
## mlogit(formula = sp_age_diff_cat ~ 1 | CENGENDR, data = hhrel_mlogit_data,
##       method = "nr", print.level = 0)
##
## Frequencies of alternatives:
##      (-1,0]   (-10,-5]   (-15,-10]   (-2,-1]   (-5,-2]   (-999,-15]
##      0.0518    0.1765    0.0690    0.0437    0.1559    0.0335
##      (0,1]    (1,2]    (10,15]   (15,999]   (2,5]    (5,10]
##      0.0437    0.0577    0.0486    0.0255    0.1564    0.1376
##
## nr method
## 8 iterations, 0h:0m:2s
## g'(-H)^-lg = 3.65E-06
## successive fonction values within tolerance limits
##
## Coefficients :
##
##              Estimate Std. Error t-value Pr(>|t|)
## (-10,-5]:(intercept)    1.903    0.109   17.40 < 2e-16 ***
## (-15,-10]:(intercept)    0.961    0.120    8.01 1.1e-15 ***
## (-2,-1]:(intercept)     0.303    0.135    2.25  0.024 *
## (-5,-2]:(intercept)     1.733    0.111   15.65 < 2e-16 ***
## (-999,-15]:(intercept)   0.256    0.136    1.88  0.060 .
## (0,1]:(intercept)      -1.068    0.202   -5.29 1.2e-07 ***
## (1,2]:(intercept)      -1.520    0.241   -6.31 2.8e-10 ***
## (10,15]:(intercept)    -3.466    0.586   -5.91 3.4e-09 ***
## (15,999]:(intercept)   -4.564    1.005   -4.54 5.6e-06 ***
## (2,5]:(intercept)      -1.345    0.225   -5.99 2.1e-09 ***
## (5,10]:(intercept)     -2.485    0.368   -6.75 1.5e-11 ***
## (-10,-5]:CENGENDRmale   -3.839    0.306  -12.54 < 2e-16 ***
## (-15,-10]:CENGENDRmale  -3.744    0.437   -8.56 < 2e-16 ***
## (-2,-1]:CENGENDRmale   -1.381    0.242   -5.70 1.2e-08 ***
## (-5,-2]:CENGENDRmale   -2.670    0.221  -12.08 < 2e-16 ***
## (-999,-15]:CENGENDRmale -4.831    1.014   -4.76 1.9e-06 ***
## (0,1]:CENGENDRmale      1.361    0.242    5.62 2.0e-08 ***
## (1,2]:CENGENDRmale      2.213    0.271    8.16 2.2e-16 ***
## (10,15]:CENGENDRmale     4.073    0.600    6.79 1.1e-11 ***
## (15,999]:CENGENDRmale    4.533    1.016    4.46 8.1e-06 ***
## (2,5]:CENGENDRmale      3.095    0.250   12.38 < 2e-16 ***
## (5,10]:CENGENDRmale     4.135    0.384   10.76 < 2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Log-Likelihood: -6720
## McFadden R^2: 0.212
## Likelihood ratio test : chisq = 3620 (p.value = <2e-16)
```



```
mlogit_spouseage_odds <- cbind(odds_ratio = exp(coef(mlogit_spouseage)),
  exp(confint.default(mlogit_spouseage)))
(mlogit_spouseage_odds <- round(mlogit_spouseage_odds, 4))
```

##	odds_ratio	2.5 %	97.5 %
## (-10,-5]:(intercept)	6.7083	5.4136	8.3127
## (-15,-10]:(intercept)	2.6146	2.0666	3.3079
## (-2,-1]:(intercept)	1.3542	1.0402	1.7629
## (-5,-2]:(intercept)	5.6562	4.5529	7.0270
## (-999,-15]:(intercept)	1.2917	0.9895	1.6860
## (0,1]:(intercept)	0.3437	0.2315	0.5105
## (1,2]:(intercept)	0.2187	0.1364	0.3508
## (10,15]:(intercept)	0.0312	0.0099	0.0986
## (15,999]:(intercept)	0.0104	0.0015	0.0747
## (2,5]:(intercept)	0.2604	0.1677	0.4044
## (5,10]:(intercept)	0.0833	0.0405	0.1714
## (-10,-5]:CENGENDRmale	0.0215	0.0118	0.0392
## (-15,-10]:CENGENDRmale	0.0237	0.0100	0.0558
## (-2,-1]:CENGENDRmale	0.2512	0.1562	0.4040
## (-5,-2]:CENGENDRmale	0.0693	0.0449	0.1068
## (-999,-15]:CENGENDRmale	0.0080	0.0011	0.0583
## (0,1]:CENGENDRmale	3.8988	2.4247	6.2690
## (1,2]:CENGENDRmale	9.1429	5.3742	15.5543
## (10,15]:CENGENDRmale	58.7216	18.1262	190.2349
## (15,999]:CENGENDRmale	93.0309	12.7110	680.8861
## (2,5]:CENGENDRmale	22.0899	13.5320	36.0599
## (5,10]:CENGENDRmale	62.4742	29.4146	132.6904

```
write.csv(coef(mlogit_spouseage), file = "mlogit_spouseage_coefs.csv")
write.csv(mlogit_spouseage_odds, file = "mlogit_spouseage_odds.csv")
```