

SA2 DSC1105

Lindsay Faith Bazar and Jade Marco Morillo

May 18, 2025

Contents

0.1	Introduction	1
0.2	Variable Selection	2
0.3	Data Cleaning	4
0.4	Statistical Analyses	4

0.1 Introduction

This study focuses on identifying social and demographic predictors of pregnancy outcomes among women in the United States. The data used in this study come from the 2022–2023 Female Respondent Public Use File of the National Survey of Family Growth (NSFG), a nationally representative survey conducted by the National Center for Health Statistics (NCHS). The survey collects detailed information about family life, reproductive health, marriage, contraception, and related topics from women aged 15–49 in the U.S.

The primary objective of this analysis is to examine how a woman’s background, education, marital status, and family structure relate to the total number of pregnancies a woman have. Using Poisson and negative binomial regression models, this study explores the associations between these predictors and pregnancy numbers. Additional statistical methods, including ordinal and multinomial logistic regression, chi-square tests, and data visualizations, are used to support a deeper understanding of patterns in the data.

```
data <- read_csv("NSFG_2022_2023_FemRespPUFData.csv")
```

```
## Rows: 5586 Columns: 1912
## -- Column specification -----
## Delimiter: ","
## chr   (1): DEVICE_TYPE
## dbl (1817): CaseID, RSCRAGE, RSCRNINF, RSCRHISP, RSCRRAE, FTFMODE, AGE_R, A...
## lgl   (94): TRYADOPT_02, TRYADOPT_03, BIONUMHX_4, MARENDHX_4, ENDMARRX_Y_4, ...
##
## 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.
```

```
data
```

```
## # A tibble: 5,586 x 1,912
##   CaseID RSCRAGE RSCRNINF RSCRHISP RSCRRAE FTFMODE DEVICE_TYPE AGE_R AGESCRN
##   <dbl>   <dbl>   <dbl>   <dbl>   <dbl>   <dbl> <chr>       <dbl>   <dbl>
```

```
## 1 96064      29      5      5      3      2 Mobile      29      29
## 2 96066      18      5      1      4      2 PC          18      18
## 3 96068      37      1      5      2      2 Mobile      37      37
## 4 96071      40      1      5      3      2 PC          40      40
## 5 96072      49      1      5      2      2 PC          49      49
## 6 96074      30      1      5      2      2 Mobile      30      30
## 7 96075      25      5      5      2      2 Mobile      25      25
## 8 96080      37      1      5      3      2 Mobile      37      37
## 9 96081      44      5      5      3      2 Mobile      44      44
## 10 96082      44      1      5      3      2 Mobile      44      44
## # i 5,576 more rows
## # i 1,903 more variables: HISP <dbl>, HISPGRP <dbl>, ROSCNT <dbl>,
## #   NUMCHILD <dbl>, HHKIDS18 <dbl>, NONBIOKIDS <dbl>, MARSTAT <dbl>,
## #   LMARSTAT <dbl>, RMARIT <dbl>, EVRMARRY <dbl>, SSMARCOH <dbl>, MANREL <dbl>,
## #   EARNHS_Y <dbl>, MYSCHOL_Y <dbl>, EARNBA_Y <dbl>, WTHPARNW <dbl>,
## #   ONOWN <dbl>, ONOWN18 <dbl>, INTACT <dbl>, PARMARR <dbl>, INTACT18 <dbl>,
## #   LVSIT14F <dbl>, LVSIT14M <dbl>, WOMRASDU <dbl>, MOMWORKD <dbl>, ...
```

A subset of variables was selected for this analysis:

Variable	Description
PREGNUM	Total number of pregnancies reported by the respondent (response variable).
FMARITAL	Current formal marital status of the respondent (e.g., married, divorced, never married).
HISPRACE	Respondent's race and Hispanic origin, categorized as White, Black, or Other.
HIEDUC	Highest level of education completed by the respondent, treated as an ordered categorical variable.
AGER	Age of the respondent at the time of the interview (continuous variable).
EDUCMOM	Education level of the respondent's mother or female caregiver, used as a proxy for childhood socioeconomic background.
INTCTFAM	Indicates whether the respondent grew up in an intact family (i.e., living with both biological or adoptive parents).

0.2 Variable Selection

0.2.0.1 *Dependent Variable (Y)* Count Variable

PREGNUM - Total Number of Pregnancies

This is the main outcome of the analysis. We want to understand what factors are related to how many times a woman has been pregnant.

```
summary(data$PREGNUM)
```

```
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##      0.000   0.000   1.000   1.476   2.000   16.000
```

0.2.0.2 *Independent Variable (X_i)* Categorical (nominal)

FMARITAL - Marital Status

Whether someone is married, divorced, or never married can influence their chances of getting pregnant. For example, married women might be more likely to plan for children.

```
summary(data$FMARITAL)
```

##	Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
##	1.000	1.000	5.000	3.361	5.000	5.000

HISPRACE2 - Race and Hispanic Origin

This can help see if pregnancy patterns are different across group, possibly due to things like culture or access to healthcare.

```
summary(data$FMARITAL)
```

##	Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
##	1.000	1.000	5.000	3.361	5.000	5.000

Ordered Categorical

HIEDUC - Highest Educational Level

For example, women with more education might focus on school or work first and have fewer, later, or no plans of pregnancies.

```
summary(data$HIEDUC)
```

##	Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
##	1.000	4.000	6.000	5.863	8.000	11.000

Demographics

AGER - Age

The older someone is, the more time they've had to potentially have pregnancies.

```
summary(data$AGER)
```

##	Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
##	15.00	25.00	33.00	32.36	40.00	50.00

EDUCMOM- Mother's Education

Growing up with an educated mother might shape attitudes towards family planning.

```
summary(data$EDUCMOM)
```

##	Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
##	1.000	2.000	3.000	3.926	4.000	95.000

INTCTFAM - Grew Up in an Intact Family or Not

This looks at whether women lived with both parents growing up. Family structure might affect emotional development and future life choices, including when or whether to have children.

```
summary(data$INTCTFAM)
```

```
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##      1.000   1.000   1.000   1.309   2.000   2.000
```

0.3 Data Cleaning

Checking null values:

```
vars <- c("PREGNUM", "FMARITAL", "HIEDUC", "HISPRACE2", "AGER", "EDUCMOM", "INTCTFAM")

sapply(data[vars], function(x) sum(is.na(x)))
```

```
##      PREGNUM  FMARITAL    HIEDUC HISPRACE2      AGER  EDUCMOM  INTCTFAM
##           0          0          0          0          0          0          0
```

Recoding categorical variables:

```
data <- data %>%
  mutate(
    FMARITAL = factor(FMARITAL,
                      levels = c(1, 2, 3, 4, 5),
                      labels = c("Married", "Widowed", "Divorced", "Separated", "Never Married")),
    HISPRACE2 = factor(HISPRACE2,
                      levels = c(1, 2, 3, 4),
                      labels = c("Hispanic", "White", "Black", "Other")),
    HIEDUC = factor(HIEDUC, ordered = TRUE),
    EDUCMOM = factor(EDUCMOM, ordered = TRUE),
    INTCTFAM = factor(INTCTFAM, levels = c(1, 2), labels = c("Intact", "Not Intact"))
  )
```

0.4 Statistical Analyses

```
poisson_model <- glm(PREGNUM ~ FMARITAL + HISPRACE2 + HIEDUC + AGER + EDUCMOM + INTCTFAM,
                     data = data)
```

```
summary(poisson_model)
```

0.4.0.1 Poisson Regression

```
##
## Call:
## glm(formula = PREGNUM ~ FMARITAL + HISPRACE2 + HIEDUC + AGER +
##      EDUCMOM + INTCTFAM, family = poisson(link = "log"), data = data)
##
## Coefficients:
```

```
##               Estimate Std. Error z value Pr(>|z|)
## (Intercept)    -0.892588   0.066881 -13.346 < 2e-16 ***
## FMARITALWidowed -0.262698   0.127781  -2.056 0.039797 *
## FMARITALDivorced -0.158446   0.037558  -4.219 2.46e-05 ***
## FMARITALSeparated -0.049787   0.058604  -0.850 0.395579
## FMARITALNever Married -0.783609 0.028664 -27.338 < 2e-16 ***
## HISPRACE2White   -0.061340   0.031257  -1.962 0.049709 *
## HISPRACE2Black    0.266027   0.036959   7.198 6.12e-13 ***
## HISPRACE2Other   -0.150477   0.042451  -3.545 0.000393 ***
## HIEDUC.L         -0.584296   0.065571  -8.911 < 2e-16 ***
## HIEDUC.Q         -0.145761   0.056741  -2.569 0.010202 *
## HIEDUC.C          0.320143   0.051383   6.231 4.65e-10 ***
## HIEDUC^4         -0.005446   0.054696  -0.100 0.920690
## HIEDUC^5          0.060968   0.054196   1.125 0.260601
## HIEDUC^6         -0.098124   0.051321  -1.912 0.055878 .
## HIEDUC^7         -0.037738   0.047820  -0.789 0.430015
## HIEDUC^8         -0.012116   0.040001  -0.303 0.761971
## HIEDUC^9         -0.007563   0.034637  -0.218 0.827159
## HIEDUC^10         0.025773   0.036457   0.707 0.479595
## AGER             0.045455   0.001477  30.771 < 2e-16 ***
## EDUCMOM.L        -0.117960   0.056100  -2.103 0.035496 *
## EDUCMOM.Q         0.173696   0.049688   3.496 0.000473 ***
## EDUCMOM.C         0.098009   0.034487   2.842 0.004485 **
## EDUCMOM^4         0.078696   0.024613   3.197 0.001387 **
## INTCTFAMNot Intact 0.229787   0.024145   9.517 < 2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for poisson family taken to be 1)
##
## Null deviance: 12582.0 on 5585 degrees of freedom
## Residual deviance: 8731.9 on 5562 degrees of freedom
## AIC: 17206
##
## Number of Fisher Scoring iterations: 6
```

To assess whether the Poisson regression model was appropriate, we checked for overdispersion, by examining the ratio of the residual deviance to its degrees of freedom.

```
dispersion <- sum(residuals(poisson_model, type = "pearson")^2) / poisson_model$df.residual
dispersion
```

```
## [1] 1.63383
```

The Poisson regression model showed evidence of overdispersion (dispersion statistic > 1).

To address this, a Negative Binomial model was fitted to account for the extra-Poisson variation.

```
nb_model <- glm.nb(PREGNUM ~ FMARITAL + HISPRACE2 + HIEDUC + AGER + EDUCMOM + INTCTFAM,
                   data = data)
summary(nb_model)
```

```
##
```

```

## Call:
## glm.nb(formula = PREGNUM ~ FMARITAL + HISPRACE2 + HIEDUC + AGER +
##       EDUCMOM + INTCTFAM, data = data, init.theta = 2.346770487,
##       link = log)
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)      -1.171003   0.088576 -13.220 < 2e-16 ***
## FMARITALWidowed    -0.308018   0.184903  -1.666  0.09575 .
## FMARITALDivorced   -0.243232   0.054254  -4.483  7.35e-06 ***
## FMARITALSeparated  -0.083390   0.087683  -0.951  0.34158
## FMARITALNever Married -0.876554  0.037591 -23.318 < 2e-16 ***
## HISPRACE2White     -0.062713   0.042515  -1.475  0.14019
## HISPRACE2Black      0.323478   0.050895   6.356  2.07e-10 ***
## HISPRACE2Other     -0.165591   0.056771  -2.917  0.00354 **
## HIEDUC.L           -0.686856   0.089277  -7.694  1.43e-14 ***
## HIEDUC.Q            -0.218934   0.076444  -2.864  0.00418 **
## HIEDUC.C            0.415490   0.069475   5.980  2.23e-09 ***
## HIEDUC^4            -0.028068   0.074604  -0.376  0.70675
## HIEDUC^5            0.078777   0.073555   1.071  0.28417
## HIEDUC^6            -0.088840   0.070085  -1.268  0.20493
## HIEDUC^7            -0.071344   0.065955  -1.082  0.27938
## HIEDUC^8            0.019116   0.055448   0.345  0.73028
## HIEDUC^9            -0.016349   0.047437  -0.345  0.73036
## HIEDUC^10           0.042292   0.050256   0.842  0.40006
## AGER                0.054487   0.001976  27.568 < 2e-16 ***
## EDUCMOM.L           -0.130601   0.079521  -1.642  0.10052
## EDUCMOM.Q           0.220303   0.070078   3.144  0.00167 **
## EDUCMOM.C           0.131053   0.047777   2.743  0.00609 **
## EDUCMOM^4           0.080844   0.033125   2.441  0.01466 *
## INTCTFAMNot Intact  0.259201   0.033146   7.820  5.28e-15 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for Negative Binomial(2.3468) family taken to be 1)
##
## Null deviance: 8370.1 on 5585 degrees of freedom
## Residual deviance: 5709.3 on 5562 degrees of freedom
## AIC: 16496
##
## Number of Fisher Scoring iterations: 1
##
##              Theta:  2.347
##            Std. Err.:  0.131
##
## 2 x log-likelihood: -16445.707

```

Model fit was compared using the Akaike Information Criterion (AIC).

```
AIC(poisson_model, nb_model)
```

```
##              df      AIC
```

```
## poisson_model 24 17206.42
## nb_model      25 16495.71
```

The Negative Binomial model provided a better fit to the data, with a lower AIC (X) compared to the Poisson model (Y). Based on these results, the Negative Binomial model was considered the more appropriate model for inference.

Summary of significant predictors:

```
coefs <- coef(nb_model)

IRRs <- exp(coefs)

percent_change <- (IRRs - 1) * 100

pvalues <- summary(nb_model)$coefficients[, "Pr(>|z|)"]

table <- data.frame(
  Estimate = coefs,
  IRR = IRRs,
  Percent_Change = round(percent_change, 2),
  P_Value = pvalues
)

significant_predictors <- table %>% filter(P_Value < 0.05)

significant_predictors
```

##	Estimate	IRR	Percent_Change	P_Value
## (Intercept)	-1.17100274	0.3100559	-68.99	6.699006e-40
## FMARITALDivorced	-0.24323236	0.7840893	-21.59	7.352042e-06
## FMARITALNever Married	-0.87655424	0.4162146	-58.38	2.918191e-120
## HISPRACE2Black	0.32347775	1.3819254	38.19	2.074282e-10
## HISPRACE2Other	-0.16559115	0.8473926	-15.26	3.536329e-03
## HIEDUC.L	-0.68685607	0.5031555	-49.68	1.431509e-14
## HIEDUC.Q	-0.21893403	0.8033747	-19.66	4.183537e-03
## HIEDUC.C	0.41549045	1.5151136	51.51	2.226064e-09
## AGER	0.05448669	1.0559984	5.60	2.716662e-167
## EDUCMOM.Q	0.22030261	1.2464539	24.65	1.668472e-03
## EDUCMOM.C	0.13105287	1.1400280	14.00	6.088342e-03
## EDUCMOM^4	0.08084384	1.0842016	8.42	1.466484e-02
## INTCTFAMNot Intact	0.25920087	1.2958941	29.59	5.281576e-15

The negative binomial regression model revealed several important factors associated with the number of pregnancies among women. Compared to married women, those who were never married had 58% fewer pregnancies (IRR = 0.42), and those who were divorced had 22% fewer pregnancies (IRR = 0.78). Widowed women also had fewer pregnancies (IRR = 0.73).

In terms of race, Black women had 38% more pregnancies than Hispanic women (IRR = 1.38), while women classified as Other races had 15% fewer (IRR = 0.85). White women had slightly fewer pregnancies (IRR = 0.94).

Education showed a clear pattern: as education increased, the number of pregnancies generally decreased. For example, the main trend (HIEDUC.L) showed that each step up in education level was associated with a 50% decrease in the number of pregnancies (IRR = 0.50).

Age also had a strong positive effect: for each additional year of age, the expected number of pregnancies increased by about 5.6% (IRR = 1.06), as older women naturally have had more time to become pregnant.

Looking at background factors, women whose mothers had some college education had 25% more pregnancies (IRR = 1.25), while those whose mothers had less than high school had slightly fewer (IRR = 0.88). Additionally, women who did not grow up in an intact family had 30% more pregnancies than those who did (IRR = 1.30),

0.4.0.2 Contingency Tables Distribution of Marital Status by Education Level:

```
ct <- table(na.omit(data[, c("FMARITAL", "HIEDUC"))))
ct
```

```
##           HIEDUC
## FMARITAL      1  2  3  4  5  6  7  8  9 10 11
## Married      85 30 44 164 254 147 110 637 437 90 62
## Widowed       5  0  1  1  1  5  2  6  3  0  1
## Divorced      12  4 16  45  80  34  28  80  51  9  6
## Separated     15  6  4  18  33  13  7  11  3  0  2
## Never Married 506 129 88 579 588 154 135 533 258 36 18
```

Chi-Square Test

```
chi <- chisq.test(ct)
chi
```

```
##
## Pearson's Chi-squared test
##
## data:  ct
## X-squared = 804.43, df = 40, p-value < 2.2e-16
```

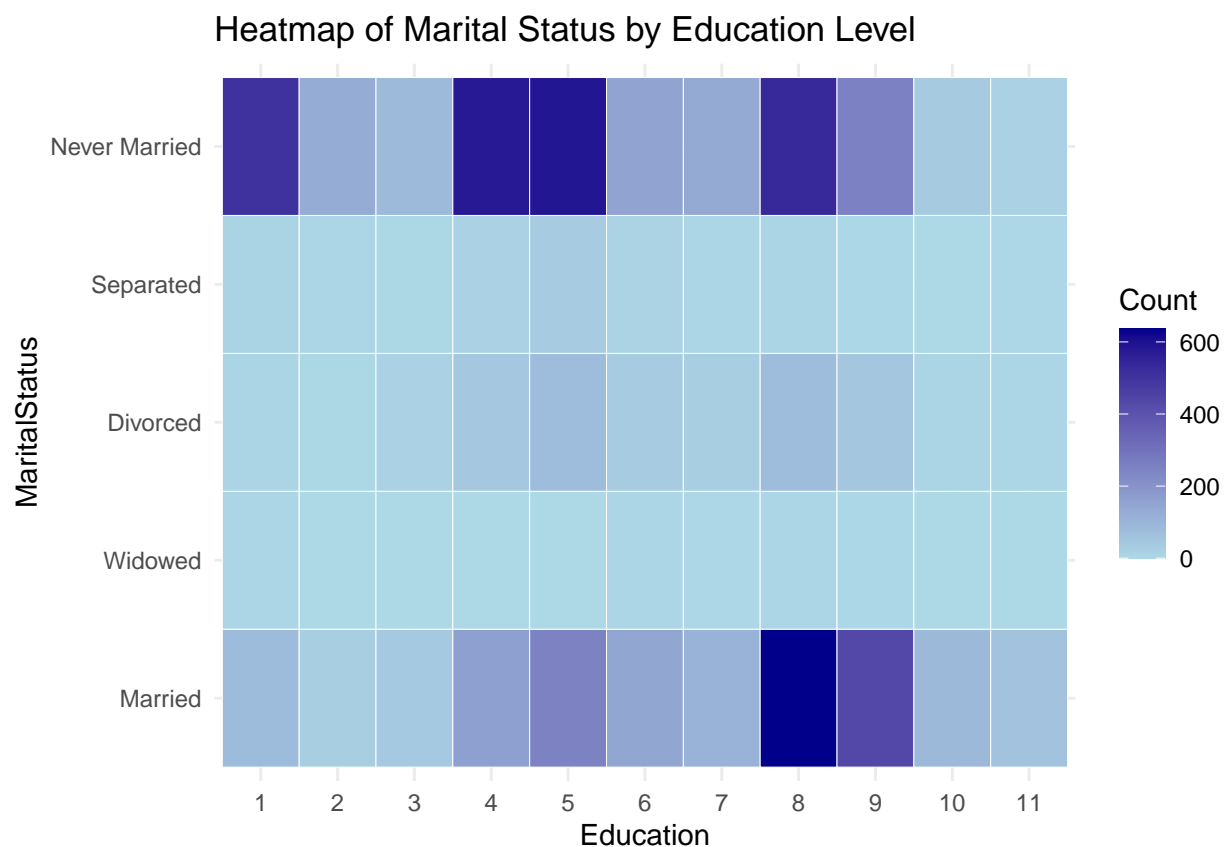
The Pearson's Chi-squared test of independence indicated a statistically significant association between marital status and highest educational level. ($p < 2.2e-16$). This suggests that the distribution of education levels differs significantly across marital status categories.

Heatmap

```
library(ggplot2)

ct_df <- as.data.frame(ct)
names(ct_df) <- c("MaritalStatus", "Education", "Count")

ggplot(ct_df, aes(x = Education, y = MaritalStatus, fill = Count)) +
  geom_tile(color = "white") +
  scale_fill_gradient(low = "lightblue", high = "darkblue") +
  labs(title = "Heatmap of Marital Status by Education Level") +
  theme_minimal()
```

0.4.0.3 Categorical Response Modeling Ordinal Logistic Regression (HIEDUC)

```
ord_model <- polr(HIEDUC ~ AGER + HISPRACE2 + INTCTFAM + EDUCMOM, data = data, Hess = TRUE)
```

```
summary(ord_model)
```

```
## Call:
## polr(formula = HIEDUC ~ AGER + HISPRACE2 + INTCTFAM + EDUCMOM,
##       data = data, Hess = TRUE)
##
## Coefficients:
##               Value Std. Error t value
## AGER              0.08940   0.00282  31.700
## HISPRACE2White     0.41712   0.06688   6.237
## HISPRACE2Black    -0.08701   0.08194  -1.062
## HISPRACE2Other     0.82256   0.08788   9.360
## INTCTFAMNot Intact -0.61583   0.05362 -11.486
## EDUCMOM.L          0.59627   0.14481   4.118
## EDUCMOM.Q         -0.99603   0.12586  -7.914
## EDUCMOM.C         -0.49350   0.08275  -5.963
## EDUCMOM^4         -0.23277   0.05267  -4.420
##
## Intercepts:
##      Value Std. Error t value
## 1|2    0.6613   0.1095   6.0382
```

```
## 2|3      0.9830  0.1088    9.0329
## 3|4      1.2296  0.1085   11.3299
## 4|5      2.2246  0.1101   20.2030
## 5|6      3.1571  0.1143   27.6332
## 6|7      3.4809  0.1157   30.0755
## 7|8      3.7430  0.1169   32.0222
## 8|9      5.1388  0.1242   41.3677
## 9|10     6.9035  0.1406   49.1129
## 10|11    7.8775  0.1634   48.2170
##
## Residual Deviance: 21710.97
## AIC: 21748.97
```

Odd Ratios

```
exp(coef(ord_model))
```

```
##              AGER      HISPRA2White      HISPRA2Black      HISPRA2Other
##      1.0935126      1.5175921      0.9166692      2.2763287
## INTCTFAMNot Intact      EDUCMOM.L      EDUCMOM.Q      EDUCMOM.C
##      0.5401950      1.8153335      0.3693430      0.6104877
##      EDUCMOM^4
##      0.7923370
```

An ordinal logistic regression revealed that age was positively associated with higher educational attainment (OR = 1.09, $p < .001$). Compared to the hispanics, White individuals (OR = 1.52, $p < .001$) and those of Other races (OR = 2.28, $p < .001$) had higher odds of achieving higher education levels. Coming from a non-intact family significantly reduced the odds of attaining higher education (OR = 0.54, $p < .001$). While maternal education showed a complex relationship, with both significant positive and negative coefficients.

Multinomial Logistic Regression (FMARITAL)

```
library(nnet)
```

```
multi_model <- multinom(FMARITAL ~ AGER + HIEDUC + HISPRA2 + INTCTFAM + EDUCMOM, data = data)
```

```
## # weights: 105 (80 variable)
## initial value 8990.320179
## iter 10 value 4788.946750
## iter 20 value 4570.514435
## iter 30 value 4329.546878
## iter 40 value 4307.172512
## iter 50 value 4294.564173
## iter 60 value 4288.023850
## iter 70 value 4285.055984
## iter 80 value 4284.876350
## iter 90 value 4284.793058
## final value 4284.785665
## converged
```

```
summary(multi_model)
```

```
## Call:
## multinom(formula = FMARITAL ~ AGER + HIEDUC + HISPRACE2 + INTCTFAM +
##      EDUCMOM, data = data)
##
## Coefficients:
##      (Intercept)          AGER  HIEDUC.L    HIEDUC.Q    HIEDUC.C
## Widowed        -13.847419  0.137375781 -1.137889 -3.73196753 -0.560036481
## Divorced        -3.876427  0.050830428 -0.673565 -0.85348318  0.508101102
## Separated       -3.910267  0.005557382 -6.890322 -2.37233086  2.144735718
## Never Married   4.250342 -0.122303787 -2.107126 -0.06123458  0.005483283
##      HIEDUC^4    HIEDUC^5    HIEDUC^6    HIEDUC^7    HIEDUC^8
## Widowed      8.84971832  0.79895015 10.0186257 -0.9243043  5.03307640
## Divorced      0.24158043 -0.25925148  0.2446356 -0.2483576  0.12511021
## Separated     5.50734340  5.87235209  4.7724076  3.7886524  1.88801847
## Never Married -0.08568934 -0.03702391  0.1292377  0.1166894 -0.04385163
##      HIEDUC^9    HIEDUC^10 HISPRACE2White HISPRACE2Black
## Widowed      0.13089716 -0.2711864    -0.3621942    1.2786401
## Divorced      0.01192105  0.2589399    0.1653316    0.6420555
## Separated     0.97614769  0.4730555    -0.8693778    0.7690865
## Never Married  0.03867706  0.1845408    -0.2263038    1.4364641
##      HISPRACE2Other INTCTFAMNot Intact  EDUCMOM.L  EDUCMOM.Q
## Widowed      0.30752855    -0.03306627 -5.5094246 -5.08976607
## Divorced     -0.44821199    0.27748924  0.1398526 -0.04459118
## Separated    -0.30105546    0.39735786  0.3038083  0.16505446
## Never Married  0.08609194    0.24118677  0.3141166 -0.25129469
##      EDUCMOM.C  EDUCMOM^4
## Widowed     -3.16924676 -1.10716310
## Divorced      0.27030043  0.19048699
## Separated     0.33063935  0.20121662
## Never Married  0.07179422  0.06649254
##
## Std. Errors:
##      (Intercept)          AGER  HIEDUC.L    HIEDUC.Q    HIEDUC.C    HIEDUC^4
## Widowed      1.6533935  0.039176289  0.7593761  0.7681827  0.8031961  0.6727713
## Divorced      0.3772580  0.008358290  0.3908232  0.3281538  0.2898631  0.3310172
## Separated     0.5167518  0.012248673  0.5211894  0.4657348  0.4729608  0.4529131
## Never Married  0.1883858  0.004633564  0.2174254  0.1900274  0.1743171  0.1862932
##      HIEDUC^5    HIEDUC^6    HIEDUC^7    HIEDUC^8    HIEDUC^9    HIEDUC^10
## Widowed      0.6191645  0.5111762  0.7702012  0.7282115  0.8167772  0.7384899
## Divorced      0.3505231  0.3315506  0.2853504  0.2264372  0.1814630  0.1809561
## Separated     0.3253042  0.3561902  0.4391397  0.4089524  0.3433274  0.3061782
## Never Married  0.1807431  0.1736410  0.1639437  0.1388889  0.1136678  0.1193634
##      HISPRACE2White HISPRACE2Black HISPRACE2Other INTCTFAMNot Intact
## Widowed      0.61882097    0.6517044    0.7155492    0.47028848
## Divorced      0.17079271    0.2269126    0.2514625    0.13044949
## Separated     0.27071404    0.2931462    0.3577169    0.20950772
## Never Married  0.09604306    0.1355093    0.1228875    0.07935933
##      EDUCMOM.L EDUCMOM.Q EDUCMOM.C  EDUCMOM^4
## Widowed      0.4447710  0.4695173  0.4044752  0.43421156
## Divorced      0.3107417  0.2737883  0.1858717  0.12522205
## Separated     0.4427407  0.3976491  0.2953838  0.22584163
```

```
## Never Married 0.2128769 0.1861256 0.1214668 0.07773155
##
## Residual Deviance: 8569.571
## AIC: 8729.571
```

Odds-Ratios

```
coefs <- summary(multi_model)$coefficients
se <- summary(multi_model)$standard.errors

zvals <- coefs / se
pvals <- 2 * (1 - pnorm(abs(zvals)))

multi_summary <- data.frame(
  Outcome = rep(rownames(coefs), each = ncol(coefs)),
  Predictor = rep(colnames(coefs), times = nrow(coefs)),
  OR = round(exp(c(coefs)), 3),
  p = round(c(pvals), 4)
)

multi_summary
```

##	Outcome	Predictor	OR	p
## 1	Widowed	(Intercept)	0.000	0.0000
## 2	Widowed	AGER	0.021	0.0000
## 3	Widowed	HIEDUC.L	0.020	0.0000
## 4	Widowed	HIEDUC.Q	70.129	0.0000
## 5	Widowed	HIEDUC.C	1.147	0.0005
## 6	Widowed	HIEDUC^4	1.052	0.0000
## 7	Widowed	HIEDUC^5	1.006	0.6500
## 8	Widowed	HIEDUC^6	0.885	0.0000
## 9	Widowed	HIEDUC^7	0.320	0.1340
## 10	Widowed	HIEDUC^8	0.510	0.0848
## 11	Widowed	HIEDUC^9	0.001	0.0000
## 12	Widowed	HIEDUC^10	0.122	0.0000
## 13	Widowed	HISPRACE2White	0.024	0.0000
## 14	Widowed	HISPRACE2Black	0.426	0.0093
## 15	Widowed	HISPRACE2Other	0.093	0.0000
## 16	Widowed	INTCTFAMNot Intact	0.941	0.7473
## 17	Widowed	EDUCMOM.L	0.571	0.4856
## 18	Widowed	EDUCMOM.Q	1.662	0.0796
## 19	Widowed	EDUCMOM.C	8.540	0.0000
## 20	Widowed	EDUCMOM^4	1.005	0.9749
## 21	Divorced	(Intercept)	6972.425	0.0000
## 22	Divorced	AGER	1.273	0.4655
## 23	Divorced	HIEDUC.L	246.495	0.0000
## 24	Divorced	HIEDUC.Q	0.918	0.6455
## 25	Divorced	HIEDUC.C	2.223	0.1969
## 26	Divorced	HIEDUC^4	0.772	0.4595
## 27	Divorced	HIEDUC^5	355.083	0.0000
## 28	Divorced	HIEDUC^6	0.964	0.8377
## 29	Divorced	HIEDUC^7	22440.569	0.0000
## 30	Divorced	HIEDUC^8	1.277	0.4606

## 31	Divorced	HIEDUC^9	118.203	0.0000
## 32	Divorced	HIEDUC^10	1.138	0.4567
## 33	Divorced	HISPRACE2White	0.397	0.2301
## 34	Divorced	HISPRACE2Black	0.780	0.3841
## 35	Divorced	HISPRACE2Other	44.197	0.0000
## 36	Divorced	INTCTFAMNot Intact	1.124	0.4766
## 37	Divorced	EDUCMOM.L	153.404	0.0000
## 38	Divorced	EDUCMOM.Q	1.133	0.5806
## 39	Divorced	EDUCMOM.C	6.606	0.0000
## 40	Divorced	EDUCMOM^4	0.957	0.7522
## 41	Separated	(Intercept)	1.140	0.8727
## 42	Separated	AGER	1.012	0.9476
## 43	Separated	HIEDUC.L	2.654	0.0045
## 44	Separated	HIEDUC.Q	1.039	0.7337
## 45	Separated	HIEDUC.C	0.762	0.7135
## 46	Separated	HIEDUC^4	1.296	0.1524
## 47	Separated	HIEDUC^5	1.605	0.1223
## 48	Separated	HIEDUC^6	1.203	0.1221
## 49	Separated	HIEDUC^7	0.696	0.5583
## 50	Separated	HIEDUC^8	1.180	0.3330
## 51	Separated	HIEDUC^9	0.419	0.0013
## 52	Separated	HIEDUC^10	0.797	0.0185
## 53	Separated	HISPRACE2White	3.592	0.0498
## 54	Separated	HISPRACE2Black	1.900	0.0047
## 55	Separated	HISPRACE2Other	2.158	0.0087
## 56	Separated	INTCTFAMNot Intact	4.206	0.0000
## 57	Separated	EDUCMOM.L	1.360	0.6674
## 58	Separated	EDUCMOM.Q	0.639	0.0747
## 59	Separated	EDUCMOM.C	0.740	0.4000
## 60	Separated	EDUCMOM^4	1.090	0.4836
## 61	Never Married	(Intercept)	0.967	0.9439
## 62	Never Married	AGER	1.320	0.0334
## 63	Never Married	HIEDUC.L	1.488	0.0579
## 64	Never Married	HIEDUC.Q	1.273	0.0024
## 65	Never Married	HIEDUC.C	0.004	0.0000
## 66	Never Married	HIEDUC^4	1.150	0.6527
## 67	Never Married	HIEDUC^5	1.355	0.4926
## 68	Never Married	HIEDUC^6	1.369	0.1401
## 69	Never Married	HIEDUC^7	0.006	0.0000
## 70	Never Married	HIEDUC^8	0.956	0.8706
## 71	Never Married	HIEDUC^9	1.179	0.6781
## 72	Never Married	HIEDUC^10	0.778	0.1770
## 73	Never Married	HISPRACE2White	0.042	0.0000
## 74	Never Married	HISPRACE2Black	1.310	0.1459
## 75	Never Married	HISPRACE2Other	1.392	0.2630
## 76	Never Married	INTCTFAMNot Intact	1.074	0.5545
## 77	Never Married	EDUCMOM.L	0.330	0.0108
## 78	Never Married	EDUCMOM.Q	1.210	0.1282
## 79	Never Married	EDUCMOM.C	1.223	0.3729
## 80	Never Married	EDUCMOM^4	1.069	0.3923

A multinomial logistic regression was used to examine how factors like age, education (both respondent's and mother's), race/ethnicity and family background affect marital status. The results showed that older age increases the chances of being widowed — for every additional year, the odds go up by 15% (OR =

1.15, $p < .001$). Black individuals were about 3.6 times more likely to be widowed ($OR = 3.59$, $p = .009$) compared to Hispanic individuals.

When it comes to being separated, White respondents were 3.6 times more likely ($OR = 3.59$, $p = .050$) and Black respondents were 1.9 times more likely ($OR = 1.90$, $p = .005$) than Hispanics. Additionally, people from non-intact families were over 4 times more likely to be separated ($OR = 4.21$, $p < .001$).

For those never married, each year of age slightly decreased the odds ($OR = 0.88$, $p = .033$). White individuals were much less likely to be never married ($OR = 0.042$, $p < .001$) compared to Hispanics. Higher maternal education also reduced the chances of never marrying — for example, a one-unit increase in a specific measure of maternal education cut the odds by about 67% ($OR = 0.33$, $p = .011$).

Model Fit

```
library(pscl)
```

```
pR2(ord_model)
```

```
## fitting null model for pseudo-r2
```

```
##           llh           llhNull           G2           McFadden           r2ML
## -1.085549e+04 -1.183263e+04  1.954282e+03  8.258024e-02  2.952088e-01
##           r2CU
## 2.995394e-01
```

```
pR2(multi_model)
```

```
## fitting null model for pseudo-r2
```

```
## # weights: 10 (4 variable)
## initial value 8990.320179
## iter 10 value 5486.601656
## final value 5479.604946
## converged
```

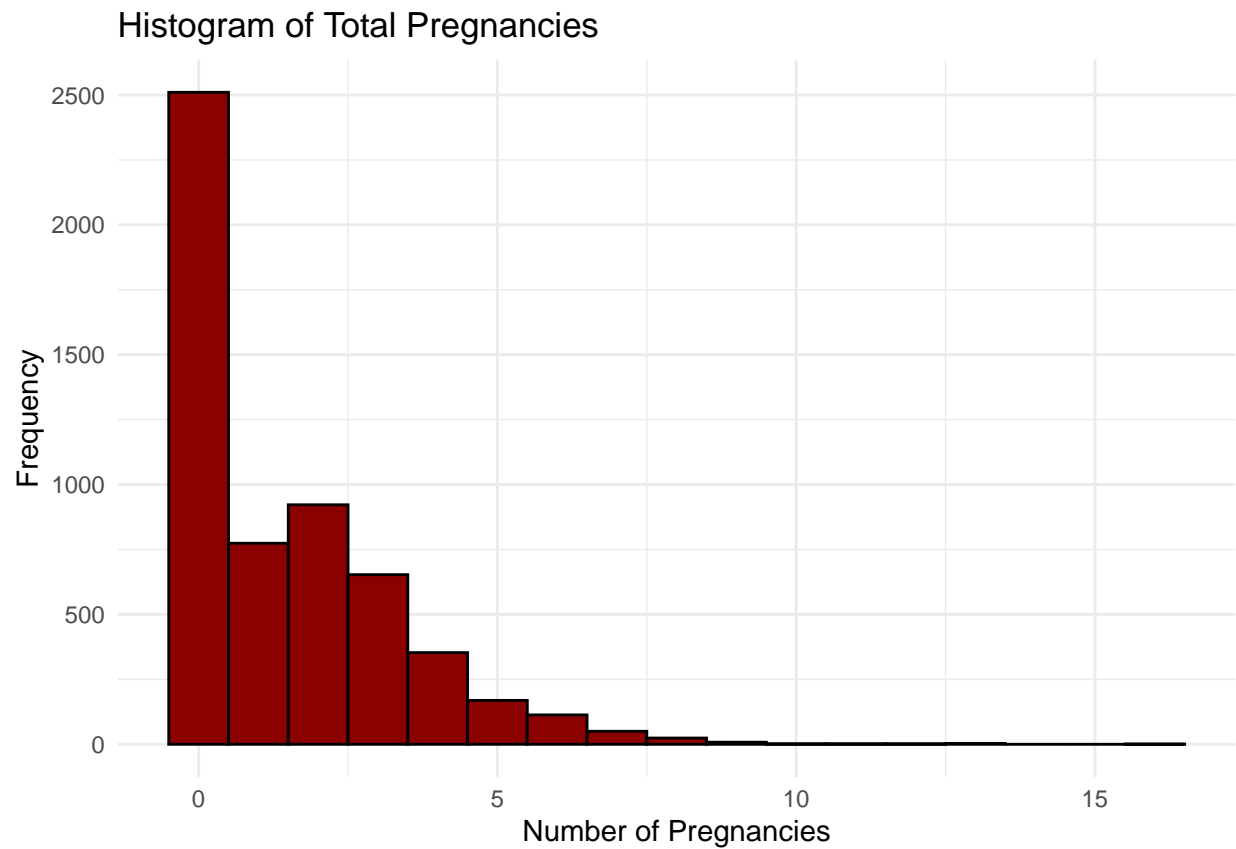
```
##           llh           llhNull           G2           McFadden           r2ML
## -4284.7856651 -5479.6049458  2389.6385614  0.2180484  0.3480521
##           r2CU
## 0.4049897
```

We assessed the model fit for both the ordinal and multinomial logistic regression models using pseudo- R^2 statistics. For the ordinal logistic regression model, the McFadden's R^2 was 0.083. The model explained approximately 29.5% (r2ML) to 30.0% (r2CU) of the variation in the outcome.

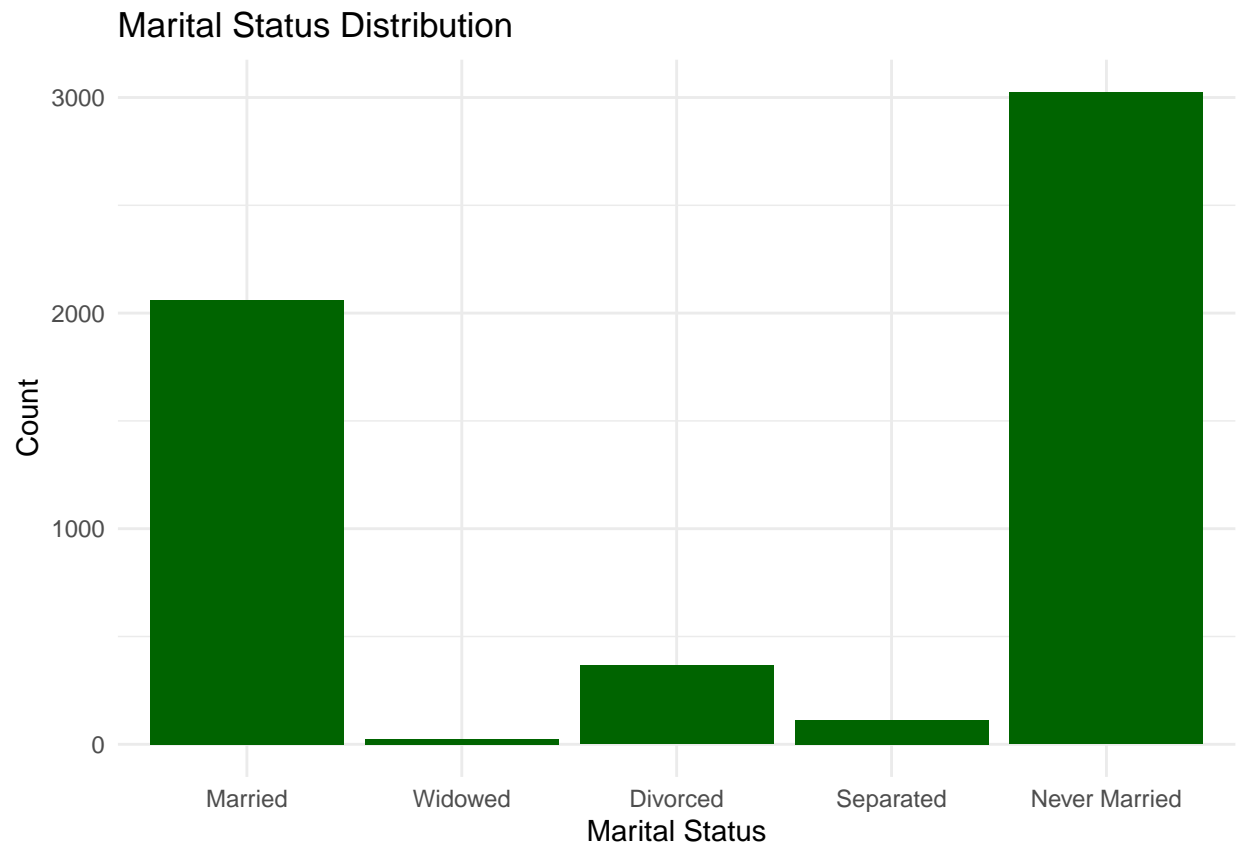
In comparison, the multinomial logistic regression model showed a better fit, with a McFadden's R^2 of 0.218. This model explained about 34.8% (r2ML) to 40.5% (r2CU) of the variance in the outcome.

0.4.0.4 EDA and Multiple Comparison Charts

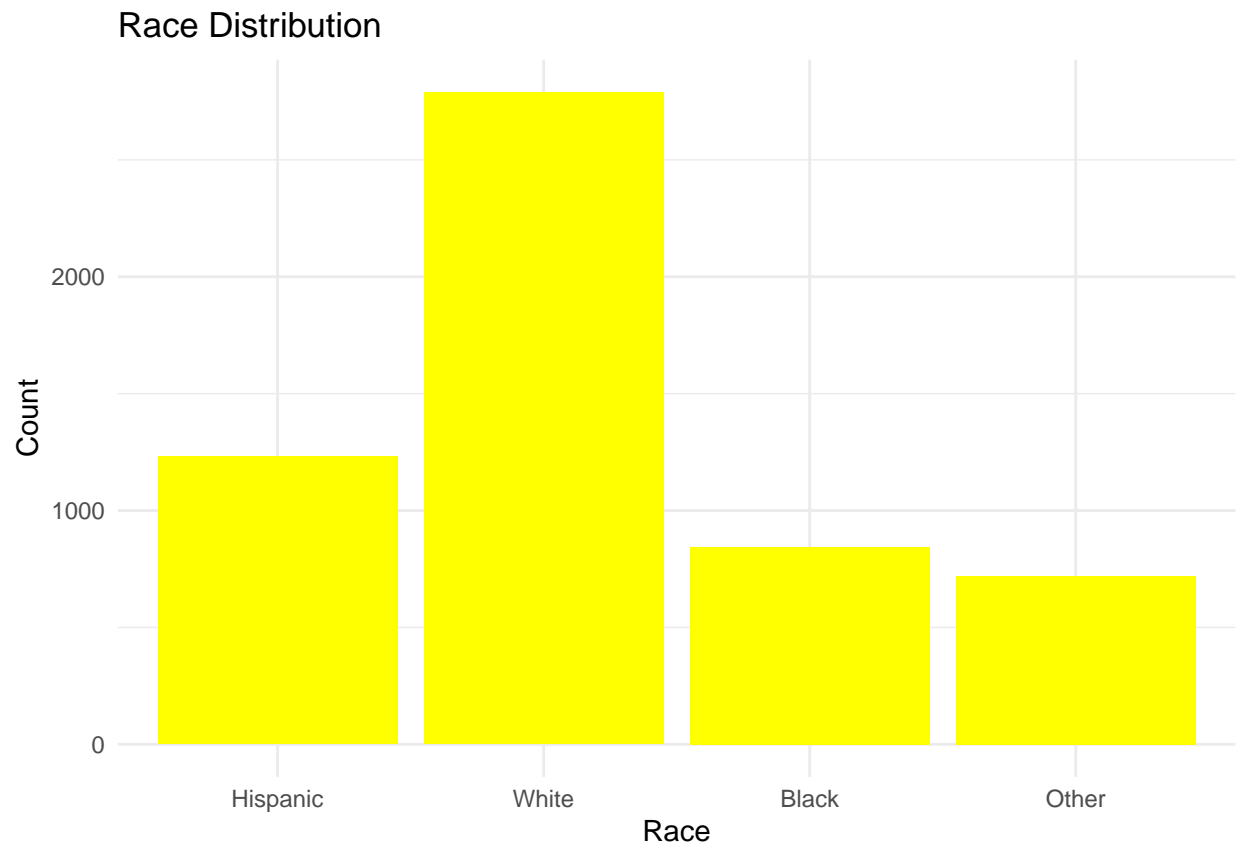
```
ggplot(data, aes(x = PREGNUM)) +
  geom_histogram(binwidth = 1, fill = "darkred", color = "black") +
  labs(title = "Histogram of Total Pregnancies", x = "Number of Pregnancies", y = "Frequency") +
  theme_minimal()
```



```
ggplot(data, aes(x = FMARITAL)) +  
  geom_bar(fill = "darkgreen") +  
  labs(title = "Marital Status Distribution", x = "Marital Status", y = "Count") +  
  theme_minimal()
```

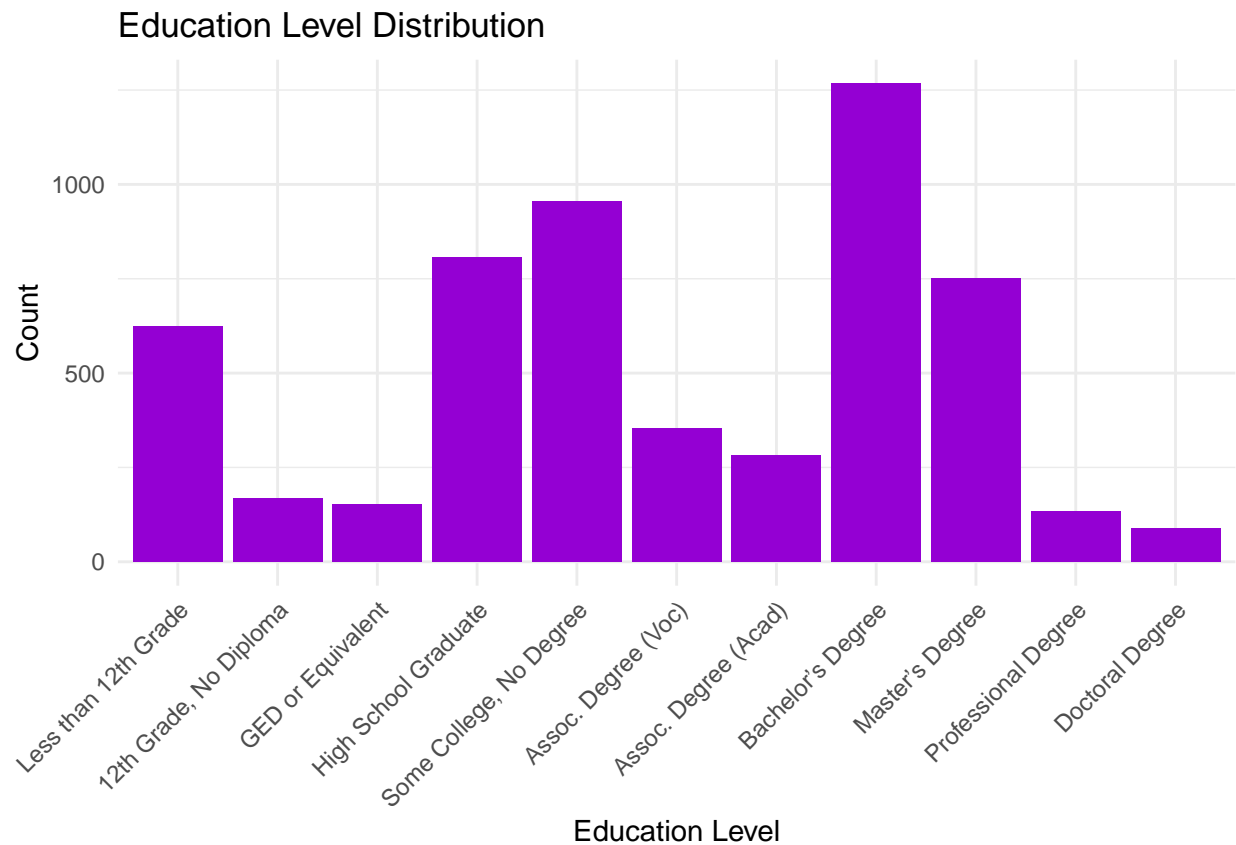


```
ggplot(data, aes(x = HISPRACE2)) +  
  geom_bar(fill = "blue") +  
  labs(title = "Race Distribution", x = "Race", y = "Count") +  
  theme_minimal()
```

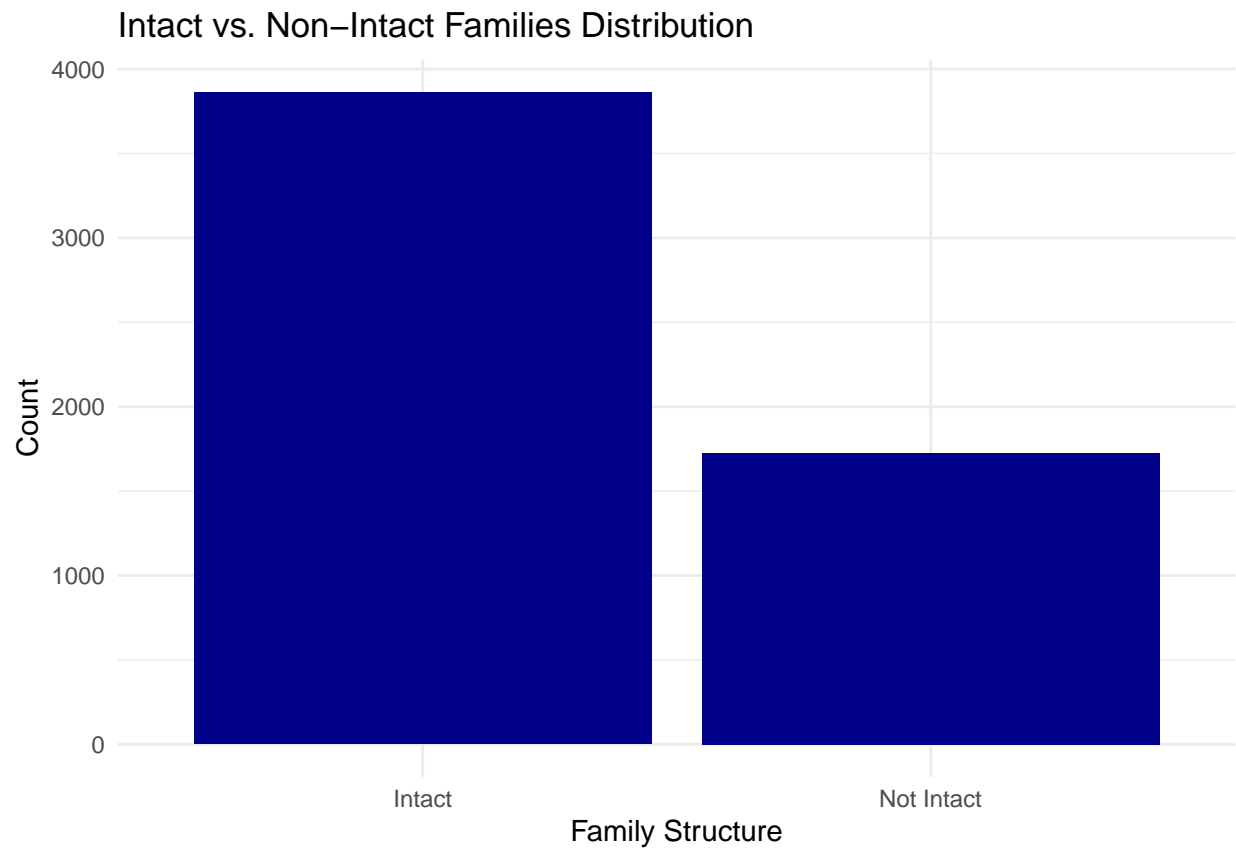



```
data$HIEDUC_f <- factor(data$HIEDUC,
  levels = 1:11,
  labels = c("Less than 12th Grade",
    "12th Grade, No Diploma",
    "GED or Equivalent",
    "High School Graduate",
    "Some College, No Degree",
    "Assoc. Degree (Voc)",
    "Assoc. Degree (Acad)",
    "Bachelor's Degree",
    "Master's Degree",
    "Professional Degree",
    "Doctoral Degree"))

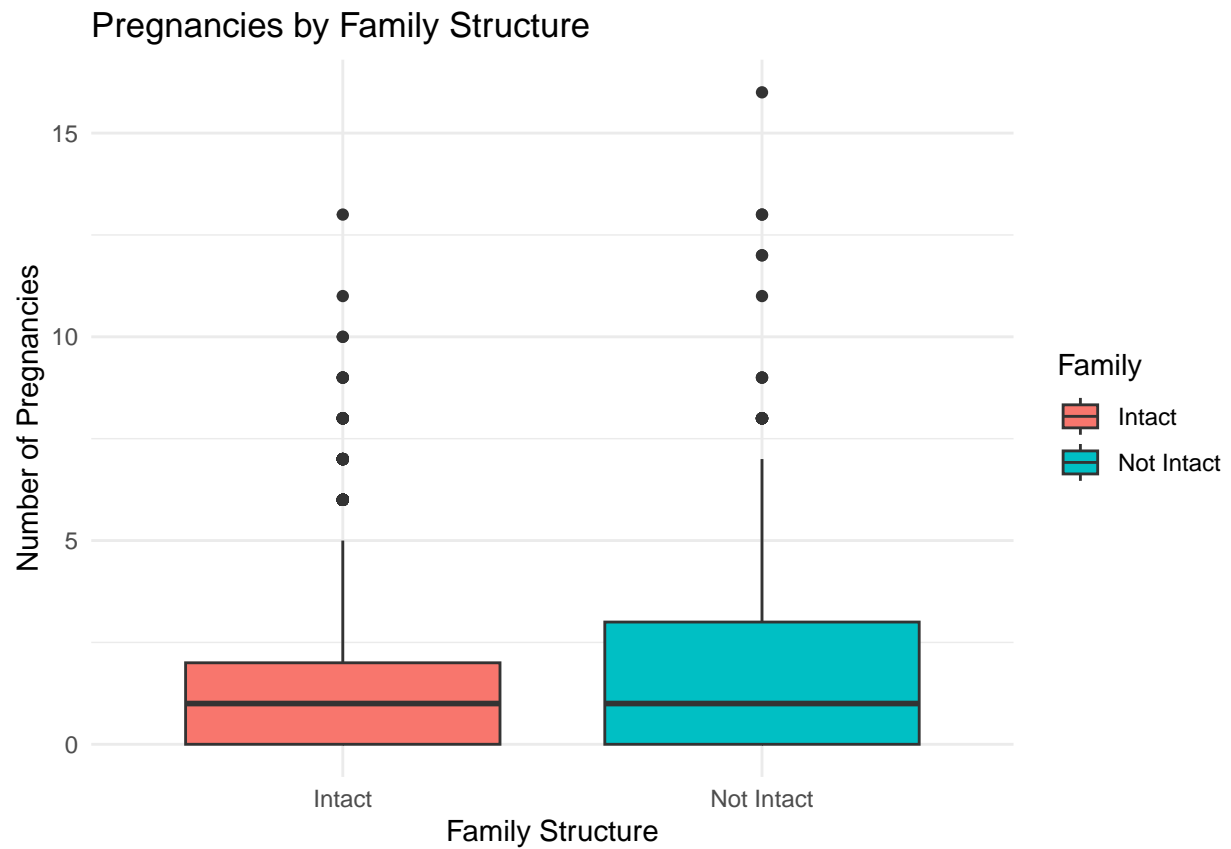
ggplot(data, aes(x = HIEDUC_f)) +
  geom_bar(fill = "darkviolet") +
  labs(title = "Education Level Distribution",
    x = "Education Level",
    y = "Count") +
  theme_minimal() +
  theme(axis.text.x = element_text(angle = 45, hjust = 1))
```



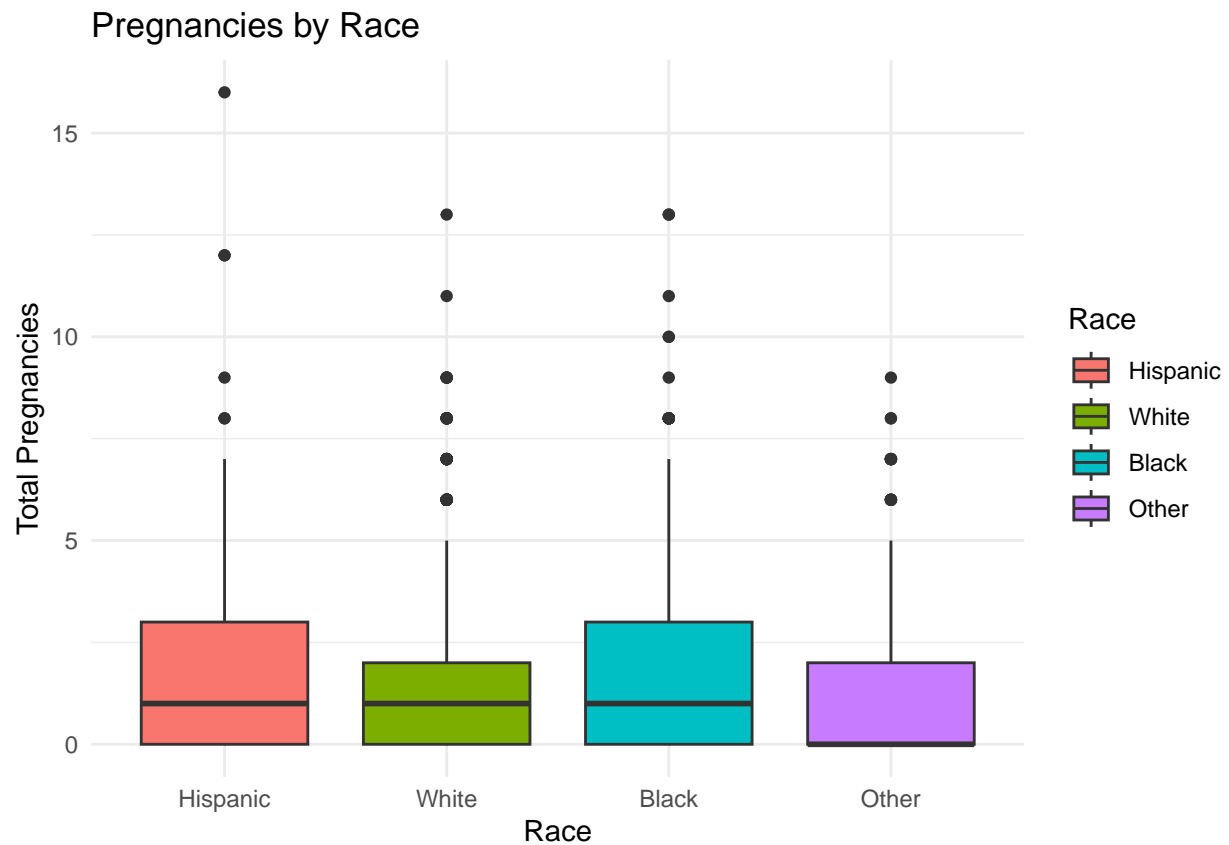
```
ggplot(data, aes(x = INTCTFAM)) +  
  geom_bar(fill="darkblue") +  
  labs(title = "Intact vs. Non-Intact Families Distribution",  
        x = "Family Structure",  
        y = "Count") +  
  theme_minimal()
```



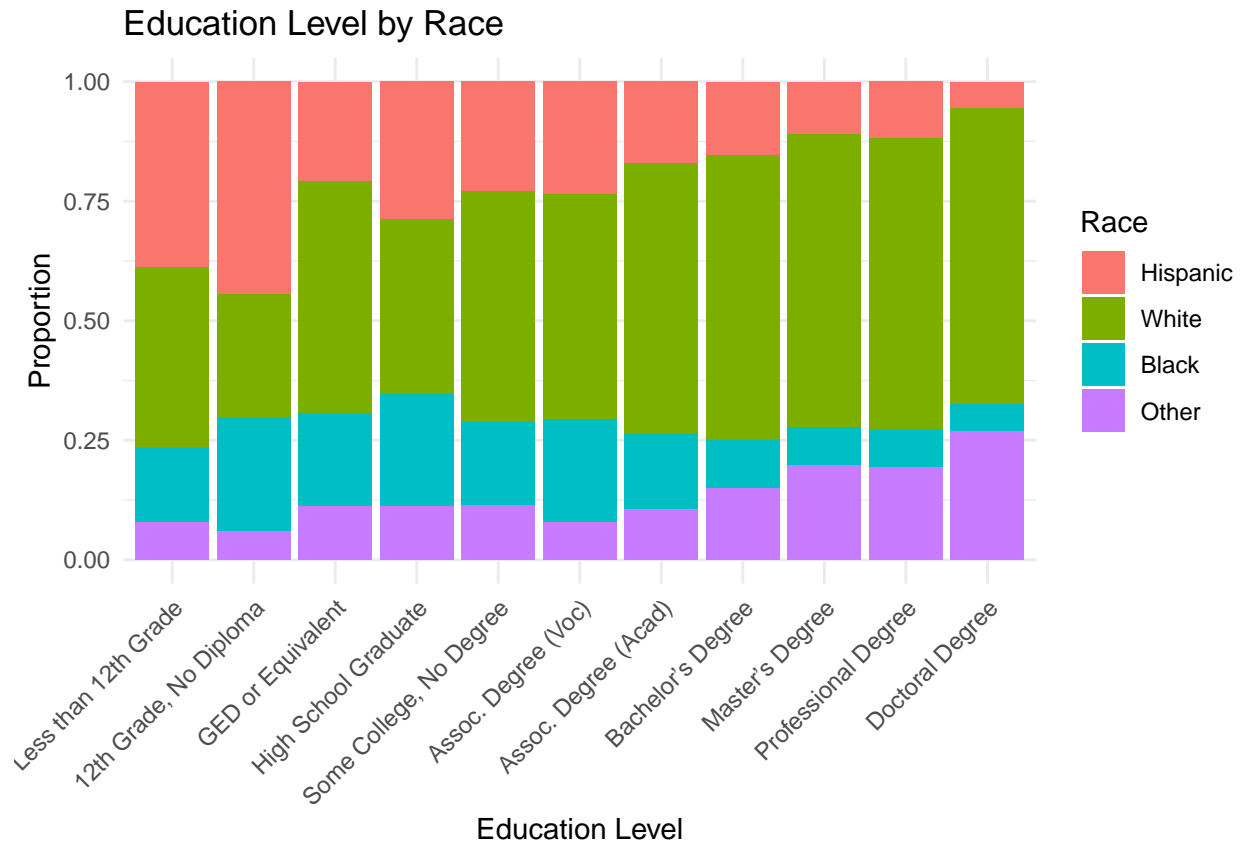
```
ggplot(data, aes(x = INTCTFAM, y = PREGNUM, fill = INTCTFAM)) +  
  geom_boxplot() +  
  labs(title = "Pregnancies by Family Structure",  
        x = "Family Structure",  
        y = "Number of Pregnancies",  
        fill = "Family") +  
  theme_minimal()
```



```
ggplot(data, aes(x = HISPRACE2, y = PREGNUM, fill = HISPRACE2)) +
  geom_boxplot() +
  labs(title = "Pregnancies by Race", x = "Race", y = "Total Pregnancies", fill = "Race") +
  theme_minimal()
```



```
ggplot(data, aes(x = HIEDUC_f, fill = HISPRACE2)) +
  geom_bar(position = "fill") +
  labs(title = "Education Level by Race",
       x = "Education Level",
       y = "Proportion",
       fill = "Race") +
  theme_minimal() +
  theme(axis.text.x = element_text(angle = 45, hjust = 1))
```



Multiple Tests

```
kruskal_test <- kruskal.test(PREGNUM ~ HISPRACE2, data = data)
kruskal_test
```

```
##
##  Kruskal-Wallis rank sum test
##
## data:  PREGNUM by HISPRACE2
## Kruskal-Wallis chi-squared = 38.536, df = 3, p-value = 2.177e-08
```

```
pairwise_result_fdr <- pairwise.wilcox.test(data$PREGNUM, data$HISPRACE2, p.adjust.method = "fdr")
pairwise_result_fdr
```

```
##
##  Pairwise comparisons using Wilcoxon rank sum test with continuity correction
##
## data:  data$PREGNUM and data$HISPRACE2
##
##      Hispanic White   Black
## White 0.05785  -      -
## Black 0.01322 1.4e-05 -
## Other 0.00022 0.00563 4.2e-08
##
## P value adjustment method: fdr
```

We compared the total number of pregnancies among different race groups using a Kruskal-Wallis test, which showed a significant difference ($\chi^2 = 38.54$, $p < 0.001$). To find out which of these groups differed, we did pairwise Wilcoxon tests with False Discovery Rate (FDR) adjustments to reduce false positives. Significant differences were found between Hispanic and Other ($p = 0.00022$), White and Black ($p = 0.00001$), Black and Other ($p < 0.00001$), and Hispanic and Black ($p = 0.01322$).