

730 Group Project

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Amani's Model: weighted linear regression

```
newdata <- read_csv("FreqCategories.csv") %>% mutate(Weight = Freq / sum(Freq))
```

```
## New names:
## Rows: 5462 Columns: 9
## -- Column specification
## ----- Delimiter: "," chr
## (2): AgeCat, EduCat dbl (7): ...1, y, REGION, SEX, RACENEW, POORYN, Freq
## 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.
## * '' -> '...1'
```

```
newdata<-mutate(newdata, weight.var=1/Freq) %>% mutate(REGION=as.factor(REGION)) %>% mutate(AgeCat=as.f
#converting y's into factor variable, changing range from 0-8 to 1-9 to match with model output
newdata1<-mutate(newdata, y=y+1) %>% mutate(y, factor(y, ordered=TRUE))
```

```
modA <- brm(
  y | weights(Weight) ~ (1 | REGION + AgeCat + SEX + RACENEW + EduCat + POORYN),
  family = gaussian(),
  data = newdata1,
  iter = 1000,
  chains = 4,
  cores = getOption("mc.cores", 4),
  seed = 12345
)
```

```
## Compiling Stan program...
## Start sampling
```

```
summary(modA)
```

```
## Family: gaussian
## Links: mu = identity; sigma = identity
## Formula: y | weights(Weight) ~ (1 | REGION + AgeCat + SEX + RACENEW + EduCat + POORYN)
```

```

## Data: newdata1 (Number of observations: 5462)
## Draws: 4 chains, each with iter = 1000; warmup = 500; thin = 1;
## total post-warmup draws = 2000
##
## Multilevel Hyperparameters:
## ~AgeCat (Number of levels: 3)
## Estimate Est.Error 1-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## sd(Intercept) 2.46 2.35 0.08 8.39 1.00 2338 992
##
## ~EduCat (Number of levels: 4)
## Estimate Est.Error 1-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## sd(Intercept) 2.39 2.12 0.09 7.93 1.00 1998 1179
##
## ~POORYN (Number of levels: 2)
## Estimate Est.Error 1-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## sd(Intercept) 2.55 2.36 0.12 8.83 1.00 1689 1088
##
## ~RACENEW (Number of levels: 6)
## Estimate Est.Error 1-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## sd(Intercept) 2.48 2.31 0.08 8.55 1.00 2080 940
##
## ~REGION (Number of levels: 4)
## Estimate Est.Error 1-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## sd(Intercept) 2.32 2.26 0.07 7.90 1.00 1739 1037
##
## ~SEX (Number of levels: 2)
## Estimate Est.Error 1-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## sd(Intercept) 2.50 2.46 0.10 8.54 1.00 2590 1320
##
## Regression Coefficients:
## Estimate Est.Error 1-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## Intercept 3.67 3.67 -3.94 10.73 1.00 2520 1185
##
## Further Distributional Parameters:
## Estimate Est.Error 1-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## sigma 5.89 4.21 1.87 14.78 1.00 2839 1446
##
## Draws were sampled using sampling(NUTS). For each parameter, Bulk_ESS
## and Tail_ESS are effective sample size measures, and Rhat is the potential
## scale reduction factor on split chains (at convergence, Rhat = 1).

```

```
prior_summary(modA)
```

```

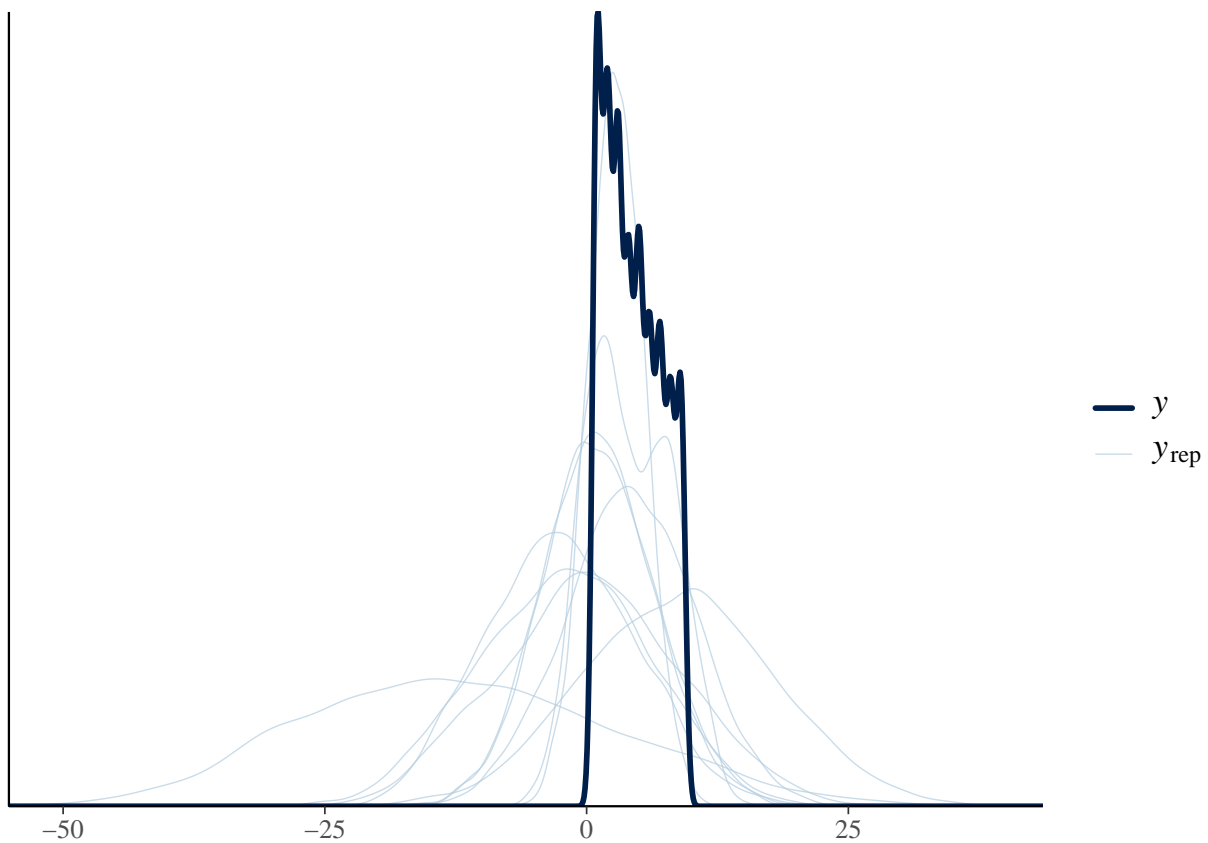
## prior class coef group resp dpar nlpar lb ub
## student_t(3, 4, 3) Intercept
## student_t(3, 0, 3) sd 0
## student_t(3, 0, 3) sd AgeCat 0
## student_t(3, 0, 3) sd Intercept AgeCat 0
## student_t(3, 0, 3) sd EduCat 0
## student_t(3, 0, 3) sd Intercept EduCat 0
## student_t(3, 0, 3) sd POORYN 0
## student_t(3, 0, 3) sd Intercept POORYN 0
## student_t(3, 0, 3) sd RACENEW 0
## student_t(3, 0, 3) sd Intercept RACENEW 0

```

```
## student_t(3, 0, 3)      sd      REGION      0
## student_t(3, 0, 3)      sd Intercept REGION      0
## student_t(3, 0, 3)      sd      SEX      0
## student_t(3, 0, 3)      sd Intercept  SEX      0
## student_t(3, 0, 3)      sigma      0
##      source
##      default
##      default
## (vectorized)
## (vectorized)
## (vectorized)
## (vectorized)
## (vectorized)
## (vectorized)
## (vectorized)
## (vectorized)
## (vectorized)
## (vectorized)
## (vectorized)
## (vectorized)
## (vectorized)
## (vectorized)
## (vectorized)
##      default
```

```
pp_check(modA)
```

```
## Using 10 posterior draws for ppc type 'dens_overlay' by default.
```



Analysis with Amani's Model

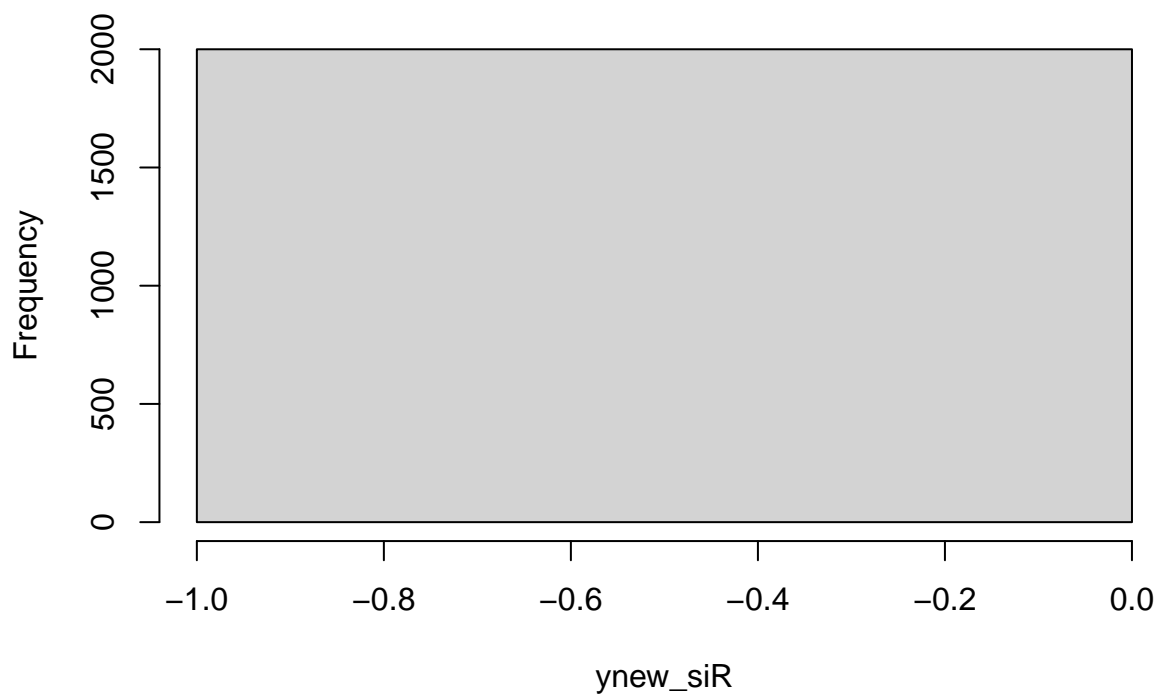
```
observed_counts <- select(newdata1, c(y, Freq))
total_freq<-group_by(observed_counts, y) %>% summarise(total=sum(Freq))
observed_props<-mutate(total_freq, observed=total/sum(total)) %>% mutate(y=as.factor(y))

get_sum_stat<-function(y, row){(sum(y==5))/nrow(row)}

tobs<-observed_props[5,3]

predicted_catsR<-as.data.frame(posterior_predict(modA))
ynew_siR<-apply(predicted_catsR, 1, get_sum_stat, newdata)
#ppc for proportion of observations in category 5
hist(ynew_siR)
abline(v = tobs)
```

Histogram of ynew_siR



```
#ppc for all categories
#formatting for ggplot
posterior_preds_longR <- predicted_catsR %>%
  pivot_longer(cols = everything(), names_to = "chain", values_to = "predicted_category")

posterior_preds_longR$predicted_category <- as.factor(posterior_preds_longR$predicted_category)
```

```

category_countsR <- table(posterior_preds_longR$predicted_category)
category_counts_dfR <- as.data.frame(category_countsR)
colnames(category_counts_dfR) <- c("y", "Count")
category_counts_propR <- mutate(category_counts_dfR, predicted=Count/(4000*5462))

combinedR <- left_join(observed_props, category_counts_propR, by="y")
combined1R <- pivot_longer(combinedR, c(3,5), names_to = "Freq")

#plot of proportion of each category for observed and predicted data
ggplot(combined1R, mapping=aes(x=y, y=value, fill=Freq))+
  geom_bar(stat="identity", position="dodge")+
  labs(title = "Mental Health Category Proportions for Observed and Predicted Data",
       x = "Category",
       y = "Proportion") +
  theme_minimal() +
  theme(axis.text.x = element_text(angle = 45, hjust = 1))

```

```

## Warning: Removed 9 rows containing missing values or values outside the scale range
## ('geom_bar()').

```

