

# 730 Group Project

Rebekah Kristal with collaborators Amani Chehimi & Shane Fitzgerald

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## Rebekah's model: weighted ordinal regression w interactions

```
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))
modR<-brm(y|weights(Freq)~REGION + AgeCat + SEX + RACENEW + EduCat + POORYN + REGION*POORYN + REGION*RA
      chains = 4,
      iter = 2000, thin = 1)
```

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

```
##
## SAMPLING FOR MODEL 'anon_model' NOW (CHAIN 1).
## Chain 1:
## Chain 1: Gradient evaluation took 0.006277 seconds
## Chain 1: 1000 transitions using 10 leapfrog steps per transition would take 62.77 seconds.
## Chain 1: Adjust your expectations accordingly!
## Chain 1:
## Chain 1:
## Chain 1: Iteration:    1 / 2000 [ 0%] (Warmup)
## Chain 1: Iteration:   200 / 2000 [10%] (Warmup)
## Chain 1: Iteration:   400 / 2000 [20%] (Warmup)
## Chain 1: Iteration:   600 / 2000 [30%] (Warmup)
## Chain 1: Iteration:   800 / 2000 [40%] (Warmup)
## Chain 1: Iteration:  1000 / 2000 [50%] (Warmup)
```

```

## Chain 1: Iteration: 1001 / 2000 [ 50%] (Sampling)
## Chain 1: Iteration: 1200 / 2000 [ 60%] (Sampling)
## Chain 1: Iteration: 1400 / 2000 [ 70%] (Sampling)
## Chain 1: Iteration: 1600 / 2000 [ 80%] (Sampling)
## Chain 1: Iteration: 1800 / 2000 [ 90%] (Sampling)
## Chain 1: Iteration: 2000 / 2000 [100%] (Sampling)
## Chain 1:
## Chain 1: Elapsed Time: 341.321 seconds (Warm-up)
## Chain 1: 272.521 seconds (Sampling)
## Chain 1: 613.842 seconds (Total)
## Chain 1:
##
## SAMPLING FOR MODEL 'anon_model' NOW (CHAIN 2).
## Chain 2:
## Chain 2: Gradient evaluation took 0.005025 seconds
## Chain 2: 1000 transitions using 10 leapfrog steps per transition would take 50.25 seconds.
## Chain 2: Adjust your expectations accordingly!
## Chain 2:
## Chain 2:
## Chain 2: Iteration: 1 / 2000 [ 0%] (Warmup)
## Chain 2: Iteration: 200 / 2000 [ 10%] (Warmup)
## Chain 2: Iteration: 400 / 2000 [ 20%] (Warmup)
## Chain 2: Iteration: 600 / 2000 [ 30%] (Warmup)
## Chain 2: Iteration: 800 / 2000 [ 40%] (Warmup)
## Chain 2: Iteration: 1000 / 2000 [ 50%] (Warmup)
## Chain 2: Iteration: 1001 / 2000 [ 50%] (Sampling)
## Chain 2: Iteration: 1200 / 2000 [ 60%] (Sampling)
## Chain 2: Iteration: 1400 / 2000 [ 70%] (Sampling)
## Chain 2: Iteration: 1600 / 2000 [ 80%] (Sampling)
## Chain 2: Iteration: 1800 / 2000 [ 90%] (Sampling)
## Chain 2: Iteration: 2000 / 2000 [100%] (Sampling)
## Chain 2:
## Chain 2: Elapsed Time: 364.68 seconds (Warm-up)
## Chain 2: 309.415 seconds (Sampling)
## Chain 2: 674.095 seconds (Total)
## Chain 2:
##
## SAMPLING FOR MODEL 'anon_model' NOW (CHAIN 3).
## Chain 3:
## Chain 3: Gradient evaluation took 0.00558 seconds
## Chain 3: 1000 transitions using 10 leapfrog steps per transition would take 55.8 seconds.
## Chain 3: Adjust your expectations accordingly!
## Chain 3:
## Chain 3:
## Chain 3: Iteration: 1 / 2000 [ 0%] (Warmup)
## Chain 3: Iteration: 200 / 2000 [ 10%] (Warmup)
## Chain 3: Iteration: 400 / 2000 [ 20%] (Warmup)
## Chain 3: Iteration: 600 / 2000 [ 30%] (Warmup)
## Chain 3: Iteration: 800 / 2000 [ 40%] (Warmup)
## Chain 3: Iteration: 1000 / 2000 [ 50%] (Warmup)
## Chain 3: Iteration: 1001 / 2000 [ 50%] (Sampling)
## Chain 3: Iteration: 1200 / 2000 [ 60%] (Sampling)
## Chain 3: Iteration: 1400 / 2000 [ 70%] (Sampling)
## Chain 3: Iteration: 1600 / 2000 [ 80%] (Sampling)

```

```

## Chain 3: Iteration: 1800 / 2000 [ 90%] (Sampling)
## Chain 3: Iteration: 2000 / 2000 [100%] (Sampling)
## Chain 3:
## Chain 3: Elapsed Time: 314.618 seconds (Warm-up)
## Chain 3: 275.246 seconds (Sampling)
## Chain 3: 589.864 seconds (Total)
## Chain 3:
##
## SAMPLING FOR MODEL 'anon_model' NOW (CHAIN 4).
## Chain 4:
## Chain 4: Gradient evaluation took 0.004786 seconds
## Chain 4: 1000 transitions using 10 leapfrog steps per transition would take 47.86 seconds.
## Chain 4: Adjust your expectations accordingly!
## Chain 4:
## Chain 4:
## Chain 4: Iteration: 1 / 2000 [ 0%] (Warmup)
## Chain 4: Iteration: 200 / 2000 [ 10%] (Warmup)
## Chain 4: Iteration: 400 / 2000 [ 20%] (Warmup)
## Chain 4: Iteration: 600 / 2000 [ 30%] (Warmup)
## Chain 4: Iteration: 800 / 2000 [ 40%] (Warmup)
## Chain 4: Iteration: 1000 / 2000 [ 50%] (Warmup)
## Chain 4: Iteration: 1001 / 2000 [ 50%] (Sampling)
## Chain 4: Iteration: 1200 / 2000 [ 60%] (Sampling)
## Chain 4: Iteration: 1400 / 2000 [ 70%] (Sampling)
## Chain 4: Iteration: 1600 / 2000 [ 80%] (Sampling)
## Chain 4: Iteration: 1800 / 2000 [ 90%] (Sampling)
## Chain 4: Iteration: 2000 / 2000 [100%] (Sampling)
## Chain 4:
## Chain 4: Elapsed Time: 2073.86 seconds (Warm-up)
## Chain 4: 346.819 seconds (Sampling)
## Chain 4: 2420.68 seconds (Total)
## Chain 4:

```

```
summary(modR)
```

```

## Family: cumulative
## Links: mu = logit; disc = identity
## Formula: y | weights(Freq) ~ REGION + AgeCat + SEX + RACENEW + EduCat + POORYN + REGION * POORYN + R
## Data: newdata1 (Number of observations: 5462)
## Draws: 4 chains, each with iter = 2000; warmup = 1000; thin = 1;
## total post-warmup draws = 4000
##
## Regression Coefficients:
##

```

	Estimate	Est.Error	1-95% CI	u-95% CI
## Intercept[1]	-0.37	0.02	-0.42	-0.33
## Intercept[2]	0.41	0.02	0.37	0.45
## Intercept[3]	1.13	0.02	1.09	1.18
## Intercept[4]	1.54	0.02	1.50	1.59
## Intercept[5]	2.08	0.02	2.04	2.13
## Intercept[6]	2.53	0.02	2.48	2.57
## Intercept[7]	3.05	0.02	3.00	3.10
## Intercept[8]	3.67	0.03	3.61	3.72
## REGION2	0.01	0.02	-0.02	0.05
## REGION3	-0.12	0.02	-0.15	-0.09

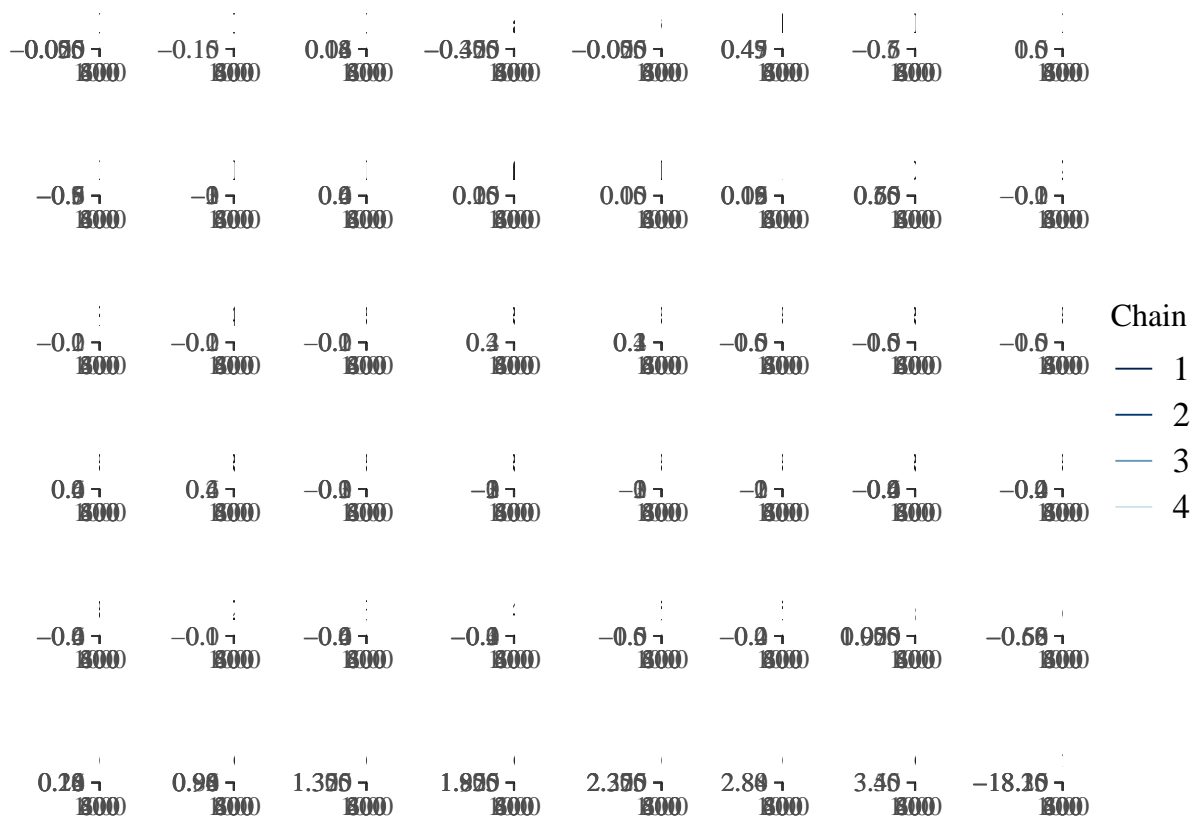
## REGION4	0.09	0.02	0.05	0.13
## AgeCatelderly	-0.38	0.01	-0.41	-0.36
## AgeCatyoungAdult	-0.03	0.01	-0.06	-0.00
## SEX2	0.45	0.01	0.43	0.47
## RACENEW200	-0.60	0.04	-0.69	-0.52
## RACENEW300	0.51	0.22	0.07	0.93
## RACENEW400	-0.69	0.06	-0.80	-0.58
## RACENEW530	-0.32	0.61	-1.52	0.83
## RACENEW541	0.35	0.11	0.13	0.57
## EduCatGraduateProfessionalorotherDegree	0.12	0.02	0.08	0.17
## EduCatHighschoolDiplomaGEDgraduate	0.02	0.02	-0.01	0.06
## EduCatSomecollegeAAorBachelorsDegree	0.12	0.02	0.09	0.16
## POORYN2	0.60	0.04	0.52	0.67
## REGION2:POORYN2	0.01	0.05	-0.08	0.11
## REGION3:POORYN2	-0.07	0.04	-0.16	0.02
## REGION4:POORYN2	-0.11	0.05	-0.21	-0.02
## REGION2:RACENEW200	0.04	0.06	-0.07	0.15
## REGION3:RACENEW200	0.30	0.05	0.21	0.40
## REGION4:RACENEW200	0.26	0.06	0.13	0.39
## REGION2:RACENEW300	-0.38	0.25	-0.87	0.11
## REGION3:RACENEW300	-0.39	0.24	-0.87	0.08
## REGION4:RACENEW300	-0.68	0.23	-1.12	-0.22
## REGION2:RACENEW400	0.27	0.08	0.11	0.44
## REGION3:RACENEW400	0.34	0.07	0.20	0.48
## REGION4:RACENEW400	0.11	0.06	-0.01	0.24
## REGION2:RACENEW530	-0.07	0.70	-1.40	1.31
## REGION3:RACENEW530	0.29	0.65	-0.97	1.58
## REGION4:RACENEW530	-0.31	0.61	-1.47	0.93
## REGION2:RACENEW541	0.08	0.14	-0.19	0.36
## REGION3:RACENEW541	0.10	0.13	-0.15	0.35
## REGION4:RACENEW541	-0.31	0.12	-0.56	-0.06
## RACENEW200:POORYN2	-0.05	0.04	-0.13	0.02
## RACENEW300:POORYN2	-0.32	0.11	-0.55	-0.11
## RACENEW400:POORYN2	-0.19	0.06	-0.31	-0.07
## RACENEW530:POORYN2	-0.23	0.28	-0.77	0.30
## RACENEW541:POORYN2	-0.07	0.09	-0.25	0.11
##	Rhat	Bulk_ESS	Tail_ESS	
## Intercept[1]	1.00	3197	2892	
## Intercept[2]	1.00	3118	2906	
## Intercept[3]	1.00	3031	2968	
## Intercept[4]	1.00	3146	2697	
## Intercept[5]	1.00	3254	2816	
## Intercept[6]	1.00	3348	2673	
## Intercept[7]	1.00	3475	2771	
## Intercept[8]	1.00	3605	3002	
## REGION2	1.00	3349	3308	
## REGION3	1.00	3613	3386	
## REGION4	1.00	3421	3307	
## AgeCatelderly	1.00	6076	2791	
## AgeCatyoungAdult	1.00	5001	2941	
## SEX2	1.00	6131	3407	
## RACENEW200	1.00	2609	2804	
## RACENEW300	1.00	2110	2528	
## RACENEW400	1.00	2396	2760	

```

## RACENEW530                1.00      2356      2613
## RACENEW541                1.00      2253      2793
## EduCatGraduateProfessionalorotherDegree 1.00      3568      2859
## EduCatHighschoolDiplomaGEDgraduate      1.00      3531      2961
## EduCatSomecollegeAAorBachelorsDegree    1.00      2778      2755
## POORYN2                      1.00      2352      2508
## REGION2:POORYN2             1.00      2927      2786
## REGION3:POORYN2             1.00      2374      2833
## REGION4:POORYN2             1.00      2816      3302
## REGION2:RACENEW200          1.00      3065      3045
## REGION3:RACENEW200          1.00      2754      2934
## REGION4:RACENEW200          1.00      3090      3121
## REGION2:RACENEW300          1.00      2248      2748
## REGION3:RACENEW300          1.00      2286      2869
## REGION4:RACENEW300          1.00      2130      2582
## REGION2:RACENEW400          1.00      3059      3232
## REGION3:RACENEW400          1.00      2925      3085
## REGION4:RACENEW400          1.00      2728      2932
## REGION2:RACENEW530          1.00      2528      2875
## REGION3:RACENEW530          1.00      2446      2916
## REGION4:RACENEW530          1.00      2388      2797
## REGION2:RACENEW541          1.00      2921      3135
## REGION3:RACENEW541          1.00      2445      2974
## REGION4:RACENEW541          1.00      2358      2635
## RACENEW200:POORYN2          1.00      5442      3033
## RACENEW300:POORYN2          1.00      4989      3266
## RACENEW400:POORYN2          1.00      5022      2978
## RACENEW530:POORYN2          1.00      4900      3009
## RACENEW541:POORYN2          1.00      5114      3113
##
## Further Distributional Parameters:
##      Estimate Est.Error l-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## disc      1.00      0.00      1.00      1.00   NA      NA      NA
##
## 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).

```

```
mcmc_trace(modR)
```



## Analysis with Rebekah's Model

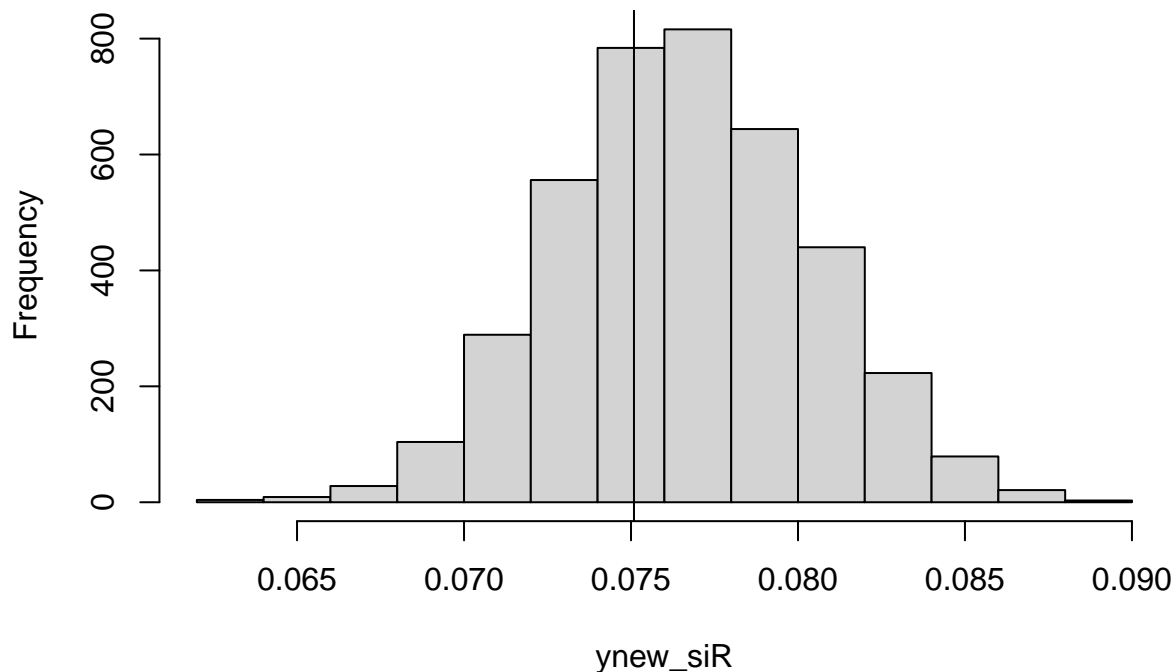
```
observed_counts <- select(newdata1, c(y, Freq))
total_freq<-group_by(observed_counts, y) %>% summarise(total=sum(Freq))
observed_props<-mutate(total_freq, observed=total/sum(total)) %>% mutate(y=as.factor(y))

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

tobs<-observed_props[5,3]

predicted_catsR<-as.data.frame(posterior_predict(modR))
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))
```

Mental Health Category Proportions for Observed and Predicted Data

