## analysis.R

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```
library(readxl)
study1 <- read_excel("Study1 Data Unrounded.xlsx")</pre>
study2 <- read_excel("Study2 Data Unrounded.xlsx")</pre>
## study 1
study1.mod1 <- glm(sent ~ trust, data = study1, family = "binomial")
summary(study1.mod1)
##
## Call:
## glm(formula = sent ~ trust, family = "binomial", data = study1)
## Deviance Residuals:
       Min
                  1Q
                        Median
                                       3Q
                                                Max
## -1.37073 -1.17373 0.06331
                                1.16488
                                            1.40913
##
## Coefficients:
             Estimate Std. Error z value Pr(>|z|)
## (Intercept) 0.9882
                           0.3708
                                   2.665 0.00770 **
                           0.1306 -2.718 0.00657 **
               -0.3550
## trust
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for binomial family taken to be 1)
##
      Null deviance: 1028.6 on 741 degrees of freedom
## Residual deviance: 1021.1 on 740 degrees of freedom
## AIC: 1025.1
##
## Number of Fisher Scoring iterations: 4
study1.mod2 <- glm(
 sent ~ trust + zAfro + attract + maturity + zfWHR + glasses + tattoos,
 data = study1, family = "binomial"
summary(study1.mod2)
##
## Call:
## glm(formula = sent ~ trust + zAfro + attract + maturity + zfWHR +
```

```
##
      glasses + tattoos, family = "binomial", data = study1)
##
## Deviance Residuals:
       Min
                  1Q
##
                        Median
                                      ЗQ
                                               Max
## -1.72997 -1.12114
                       0.02173
                               1.11971
                                           1.69625
##
## Coefficients:
              Estimate Std. Error z value Pr(>|z|)
##
## (Intercept) 2.18500
                          0.81273
                                   2.688 0.00718 **
                          0.14727 -2.778 0.00547 **
## trust
              -0.40908
## zAfro
              -0.23862
                          0.07995 -2.985 0.00284 **
                          0.14205 -1.145 0.25230
## attract
              -0.16261
## maturity
              -0.13589
                          0.08841 -1.537 0.12431
                                   3.893 9.92e-05 ***
## zfWHR
              0.32639
                          0.08385
## glasses
              0.44806
                          0.21875
                                    2.048 0.04054 *
## tattoos
              -0.55039
                          0.55835 -0.986 0.32427
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for binomial family taken to be 1)
##
##
      Null deviance: 1028.63 on 741 degrees of freedom
## Residual deviance: 986.96 on 734 degrees of freedom
## AIC: 1003
##
## Number of Fisher Scoring iterations: 4
## study 2
study2.mod1 <- glm(sent ~ trust, data = study2, family = "binomial")
summary(study2.mod1)
##
## Call:
## glm(formula = sent ~ trust, family = "binomial", data = study2)
## Deviance Residuals:
##
      Min
                1Q
                     Median
                                  3Q
                                          Max
## -2.2244 -0.8936 -0.6561
                              0.9816
                                       1.6519
##
## Coefficients:
              Estimate Std. Error z value Pr(>|z|)
## (Intercept) 5.9603
                           2.7071
                                    2.202
                                           0.0277 *
## trust
               -1.5489
                           0.6777 -2.286
                                           0.0223 *
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
      Null deviance: 51.049 on 36 degrees of freedom
## Residual deviance: 44.581 on 35 degrees of freedom
## AIC: 48.581
##
## Number of Fisher Scoring iterations: 4
```

```
study2.mod2 <- glm(</pre>
  sent ~ trust + zAfro + attract + maturity + glasses,
 data = study2, family = "binomial"
 )
summary(study2.mod2)
## Call:
## glm(formula = sent ~ trust + zAfro + attract + maturity + glasses,
      family = "binomial", data = study2)
## Deviance Residuals:
      Min 1Q Median
                                 3Q
                                         Max
## -2.4340 -0.9115 -0.5349 1.1019
                                      1.6444
##
## Coefficients:
             Estimate Std. Error z value Pr(>|z|)
## (Intercept) 7.73841
                         4.49363
                                  1.722 0.0851 .
## trust
             -1.81451
                          0.81038 -2.239
                                          0.0252 *
## zAfro
              -0.44202
                          0.38838 -1.138
                                          0.2551
## attract
              -0.47581
                          0.79120 -0.601
                                           0.5476
## maturity
              0.08854
                          0.51447
                                  0.172 0.8634
              0.72428
                          0.88392
## glasses
                                  0.819 0.4126
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for binomial family taken to be 1)
##
      Null deviance: 51.049 on 36 degrees of freedom
## Residual deviance: 42.655 on 31 degrees of freedom
## AIC: 54.655
```

## Number of Fisher Scoring iterations: 4