Appendix: Descriptive Representation and Public Support for Supreme Court Nominees

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Load the data

```
load("appendix_data.RData")

# Define variables to pass into model
data$politicized <- as.factor(data$politicized)
data$treat_age <- as.factor(data$treat_age)
data$treat_gender <- as.factor(data$treat_gender)
data$treat_race <- as.factor(data$treat_race)
data$treat_lawS <- as.factor(data$treat_lawS)
data$treat_currentP <- as.factor(data$treat_currentP)
data$treat_pAbortion <- as.factor(data$treat_pAbortion)
```

Figure A.1: Effect of Shared Descriptive Characteristics on Nominee Evaluations (5-Point Measure)

```
# Subset dataset to respondents who answered the support question
data.support <- data[!(is.na(data$support) | data$support==""), ]

# Estimate the AMCEs, cluster SEs by respondent
results.support.overall.5pt <- amce(support ~ politicized + treat_age + treat_gender + treat_race +  tr
                                    same.race, data=data.support,cluster=TRUE, respondent.id="caseid", we
summary(results.support.overall.5pt)
```

```
## ----------------------------------------
## Average Marginal Component Effects (AMCE):
## ----------------------------------------
##         Attribute Level   Estimate Std. Err  z value   Pr(>|z|)
##       politicized     1 -0.0410621 0.040625 -1.01075 3.1214e-01
##       same.gender     1  0.0298364 0.031615  0.94373 3.4531e-01
##         same.race     1  0.2071583 0.043515  4.76063 1.9299e-06 ***
##         treat_age     2  0.0075464 0.041906  0.18008 8.5709e-01
##         treat_age     3 -0.0311065 0.039372 -0.79006 4.2949e-01
##    treat_currentP     2 -0.1005992 0.048723 -2.06470 3.8952e-02   *
##    treat_currentP     3 -0.0810293 0.050543 -1.60316 1.0890e-01
##    treat_currentP     4 -0.1809825 0.048831 -3.70633 2.1029e-04 ***
##    treat_currentP     5 -0.1902778 0.050165 -3.79306 1.4880e-04 ***
##      treat_gender     2 -0.0313247 0.031684 -0.98864 3.2284e-01
##        treat_lawS     2 -0.0541058 0.044797 -1.20781 2.2712e-01
##        treat_lawS     3 -0.1059116 0.047482 -2.23056 2.5710e-02   *
##        treat_lawS     4 -0.2258504 0.044128 -5.11808 3.0867e-07 ***
##   treat_pAbortion     2 -0.2161369 0.039323 -5.49640 3.8762e-08 ***
##   treat_pAbortion     3 -0.3752037 0.050756 -7.39228 1.4433e-13 ***
```

```
##       treat_race     2  0.2432895 0.049410  4.92389 8.4841e-07 ***
##       treat_race     3  0.1538877 0.045715  3.36625 7.6198e-04 ***
## ---
## Number of Obs. = 9947
## ---
## Number of Respondents = 2499
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05
##
## --------------------
## AMCE Baseline Levels:
## --------------------
##        Attribute Level
##      politicized
##      same.gender
##        same.race
##        treat_age
##   treat_currentP
##     treat_gender
##       treat_lawS
##  treat_pAbortion
##       treat_race
```

```r
# Subset dataset to respondents who answered the qualified question
data.qualified <- data[!(is.na(data$qualified) | data$qualified==""), ]

# Estimate the AMCEs, cluster SEs by respondent
results.qualified.overall.5pt <- amce(qualified ~ politicized + treat_age + treat_gender + treat_race +
                                      same.race, data=data.qualified,cluster=TRUE, respondent.id="caseid"
summary(results.qualified.overall.5pt)
```

```
## ------------------------------------------
## Average Marginal Component Effects (AMCE):
## ------------------------------------------
##        Attribute Level   Estimate Std. Err  z value   Pr(>|z|)
##      politicized     1 -0.1158352 0.041552 -2.78775 5.3075e-03  **
##      same.gender     1  0.0090940 0.031837  0.28564 7.7515e-01
##        same.race     1  0.1615862 0.048358  3.34149 8.3332e-04 ***
##        treat_age     2  0.0075710 0.039936  0.18958 8.4964e-01
##        treat_age     3 -0.0521003 0.037224 -1.39965 1.6162e-01
##   treat_currentP     2 -0.3070378 0.052413 -5.85809 4.6822e-09 ***
##   treat_currentP     3 -0.1252704 0.052252 -2.39744 1.6510e-02   *
##   treat_currentP     4 -0.3700844 0.051205 -7.22752 4.9188e-13 ***
##   treat_currentP     5 -0.4105909 0.052945 -7.75505 8.8309e-15 ***
##     treat_gender     2  0.0033188 0.031727  0.10461 9.1669e-01
##       treat_lawS     2 -0.0921096 0.044436 -2.07285 3.8186e-02   *
##       treat_lawS     3 -0.1932003 0.048336 -3.99700 6.4151e-05 ***
##       treat_lawS     4 -0.4210037 0.045074 -9.34020 9.6154e-21 ***
##  treat_pAbortion     2 -0.1260535 0.039429 -3.19694 1.3889e-03  **
##  treat_pAbortion     3 -0.1608479 0.043091 -3.73278 1.8938e-04 ***
##       treat_race     2  0.2110381 0.048714  4.33219 1.4763e-05 ***
##       treat_race     3  0.1563189 0.050064  3.12238 1.7939e-03  **
## ---
## Number of Obs. = 9955
## ---
```

```
## Number of Respondents = 2499
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05
##
## --------------------
## AMCE Baseline Levels:
## --------------------
##        Attribute Level
##      politicized
##      same.gender
##        same.race
##        treat_age
##   treat_currentP
##     treat_gender
##       treat_lawS
##  treat_pAbortion
##       treat_race
```

```r
# Subset dataset to respondents who answered the trust question
data.trust <- data[!(is.na(data$trust) | data$trust==""), ]

# Estimate the AMCEs, cluster SEs by respondent
results.trust.overall.5pt <- amce(trust ~ politicized + treat_age + treat_gender + treat_race +  treat_
                                  same.race, data=data.trust,cluster=TRUE, respondent.id="caseid", weights
summary(results.trust.overall.5pt)
```

```
## ------------------------------------------
## Average Marginal Component Effects (AMCE):
## ------------------------------------------
##        Attribute Level   Estimate Std. Err   z value   Pr(>|z|)
##      politicized     1 -0.0117487 0.038124 -0.308169 7.5795e-01
##      same.gender     1  0.0565959 0.029180  1.939577 5.2431e-02
##        same.race     1  0.1959024 0.042717  4.586090 4.5162e-06 ***
##        treat_age     2 -0.0097536 0.035613 -0.273880 7.8418e-01
##        treat_age     3 -0.0317302 0.034636 -0.916096 3.5962e-01
##   treat_currentP     2 -0.1498465 0.047418 -3.160095 1.5772e-03  **
##   treat_currentP     3 -0.0758354 0.047235 -1.605489 1.0839e-01
##   treat_currentP     4 -0.1767371 0.046413 -3.807882 1.4016e-04 ***
##   treat_currentP     5 -0.1922830 0.044800 -4.292029 1.7705e-05 ***
##     treat_gender     2 -0.0077527 0.029113 -0.266298 7.9001e-01
##       treat_lawS     2  0.0034532 0.036717  0.094052 9.2507e-01
##       treat_lawS     3 -0.0427967 0.042808 -0.999738 3.1744e-01
##       treat_lawS     4 -0.1306075 0.041774 -3.126510 1.7689e-03  **
##  treat_pAbortion     2 -0.1349722 0.036041 -3.744962 1.8042e-04 ***
##  treat_pAbortion     3 -0.2954786 0.045143 -6.545385 5.9342e-11 ***
##       treat_race     2  0.1971656 0.045906  4.294974 1.7471e-05 ***
##       treat_race     3  0.1495151 0.044726  3.342931 8.2898e-04 ***
## ---
## Number of Obs. = 9970
## ---
## Number of Respondents = 2500
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05
##
## --------------------
```

```
## AMCE Baseline Levels:
## --------------------
##        Attribute Level
##      politicized
##      same.gender
##        same.race
##        treat_age
##   treat_currentP
##     treat_gender
##       treat_lawS
##  treat_pAbortion
##       treat_race
```

```
# Coefficients and SEs for support
# Make a coefficient plot
coef_gender_support <-0.0298364
coef_race_support <-0.2071583

se_gender_support <-0.031615
se_race_support <- 0.043515

# Create the confidence intervals
z_95 <- qnorm(.975)
ci_1 <- c(coef_gender_support - z_95*se_gender_support,
          coef_gender_support + z_95*se_gender_support)
ci_2 <- c(coef_race_support - z_95*se_race_support,
          coef_race_support + z_95*se_race_support)

gender_support <- data.frame(est = coef_gender_support,
                             lb = ci_1[1], ub = ci_1[2], model = "Support")
race_support <- data.frame(est = coef_race_support,
                             lb = ci_2[1], ub = ci_2[2], model = "Support")

# Coefficients and SEs for qualified
coef_gender_qualified <- 0.0090940
coef_race_qualified <- 0.1615862

se_gender_qualified <- 0.031837
se_race_qualified <- 0.048358

# Create the confidence intervals
z_95 <- qnorm(.975)
ci_1 <- c(coef_gender_qualified - z_95*se_gender_qualified,
          coef_gender_qualified + z_95*se_gender_qualified)
ci_2 <- c(coef_race_qualified   - z_95*se_race_qualified,
          coef_race_qualified   + z_95*se_race_qualified)

gender_qualified <- data.frame(est = coef_gender_qualified,
                             lb = ci_1[1], ub = ci_1[2], model = "Qualified")
race_qualified <- data.frame(est = coef_race_qualified,
                             lb = ci_2[1], ub = ci_2[2], model = "Qualified")

# Coefficients and SEs for trust
coef_gender_trust <-0.0565959
```

```r
coef_race_trust <- 0.1959024

se_gender_trust <-0.029180
se_race_trust <- 0.042717

# Create the confidence intervals
z_95 <- qnorm(.975)
ci_1 <- c(coef_gender_trust - z_95*se_gender_trust,
          coef_gender_trust + z_95*se_gender_trust)
ci_2 <- c(coef_race_trust   - z_95*se_race_trust,
          coef_race_trust   + z_95*se_race_trust)

gender_trust <- data.frame(est = coef_gender_trust, lb = ci_1[1],
                           ub = ci_1[2], model = "Trust")
race_trust <- data.frame(est = coef_race_trust, lb = ci_2[1],
                           ub = ci_2[2], model = "Trust")

# Make the plot
dat <- as.data.frame(c("Qualified","Trust","Support"))
colnames(dat) <- c("order")

# Race
p1 <- ggplot(race_trust, aes(x = model, y = est)) +
  geom_point() +
  geom_errorbar(aes(ymin = lb, ymax = ub), width = 0.2) +
  geom_hline(yintercept = 0, lty = 2) +
  xlab("") +
  ylab("Coefficient Estimate") +
  geom_point(data= race_qualified, aes(x = model, y = est)) +
  geom_errorbar(data=race_qualified, aes(ymin = lb, ymax = ub), width = 0.2) +
  geom_point(data= race_support, aes(x = model, y = est)) +
  geom_errorbar(data=race_support, aes(ymin = lb, ymax = ub), width = 0.2) +
  theme_bw() + coord_flip() + labs(title="Race")  + ylim(-.2,0.4) +
  scale_x_discrete(limits=dat$order) +
  theme(plot.title = element_text(hjust = 0.5))

# Gender
p2 <-ggplot(gender_trust, aes(x = model, y = est)) +
  geom_point() +
  geom_errorbar(aes(ymin = lb, ymax = ub), width = 0.2) +
  geom_hline(yintercept = 0, lty = 2) +
  xlab("") +
  ylab("Coefficient Estimate") +
  geom_point(data= gender_qualified, aes(x = model, y = est)) +
  geom_errorbar(data=gender_qualified, aes(ymin = lb, ymax = ub), width = 0.2) +
  geom_point(data= gender_support, aes(x = model, y = est)) +
  geom_errorbar(data=gender_support, aes(ymin = lb, ymax = ub), width = 0.2) +
  theme_bw() + coord_flip() + labs(title="Gender")  + ylim(-.2,0.4) +
  scale_x_discrete(limits=dat$order) +
  theme(plot.title = element_text(hjust = 0.5))

grid.arrange(p1, p2, ncol=2)
```

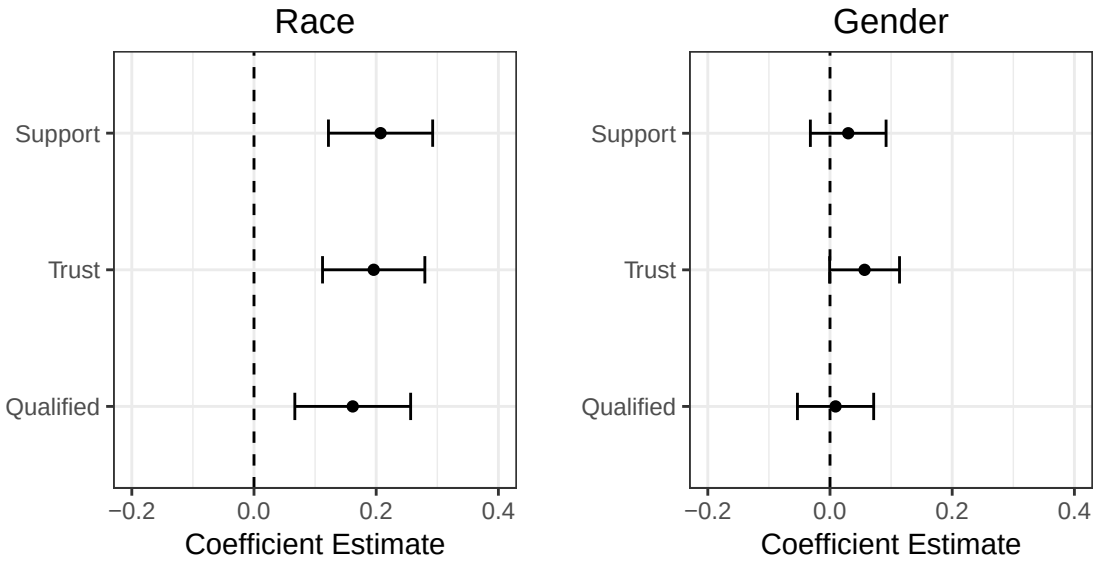


Figure A.2: Judicial Knowledge, Shared Descriptive Characteristics and Nominee Evaluations

```
# Create combined knowledge measure
data.support$know1 <- recode(data.support$Q10,"'Appointed to the bench'=1;
                             c('Elected to the bench','Neither','NA')=0")
data.support$know1 <- recode(data.support$know1,"NA=0")
data.support$know2 <- recode(data.support$Q11,"'Life term'=1;
                             c('Set number of years','Neither','NA')=0")
data.support$know2 <- recode(data.support$know2,"NA=0")
data.support$know3 <- recode(data.support$Q12,"'U.S. Supreme Court'=1;
                             c('U.S. Congress','The President','NA')=0")
data.support$know3 <- recode(data.support$know3,"NA=0")
data.support$know4 <- recode(data.support$Q13,"'John Roberts'=1;
                             c('William Rehnquist','Stephen Breyer',
                             'Antonin Scalia','Anthony Kennedy','NA')=0")
data.support$know4 <- recode(data.support$know4,"NA=0")
data.support$know5 <- recode(data.support$Q14,'"Elena Kagan"=1;
                             c("Samuel Alito","Sonia Sotomayor",
                             "Sandra Day O\'Connor","John Roberts","NA")=0')
data.support$know5 <- recode(data.support$know5,"NA=0")

data.support$judicial.knowledge <- as.numeric(as.character(data.support$know1)) +
  as.numeric(as.character(data.support$know2)) +
  as.numeric(as.character(data.support$know3)) +
  as.numeric(as.character(data.support$know4)) +
  as.numeric(as.character(data.support$know5))
data.support$judicial.knowledge.high <- as.factor(recode(data.support$judicial.knowledge,
                                                         "c(0,1,2,3)=0;c(4,5)=1") )

# Estimate the AMCEs, cluster SEs by respondent
results.support.knowledge.gender <- amce(support2 ~ politicized + treat_age + treat_gender +
                                         treat_race + same.gender*judicial.knowledge.high +
```

```
                                          same.race + treat_lawS + treat_currentP +
                                          treat_pAbortion, data=data.support, cluster=TRUE,
                                          respondent.id="caseid", weights="weight",
                                          respondent.varying="judicial.knowledge.high")
summary(results.support.knowledge.gender)
```

```
## ------------------------------------------
## Average Marginal Component Effects (AMCE):
## ------------------------------------------
##        Attribute Level    Estimate Std. Err   z value   Pr(>|z|)
##      politicized     1 -0.02152808 0.016788 -1.282321 1.9973e-01
##      same.gender     1  0.01267180 0.012395  1.022313 3.0663e-01
##        same.race     1  0.06145415 0.017882  3.436691 5.8887e-04 ***
##        treat_age     2  0.01262425 0.016684  0.756663 4.4925e-01
##        treat_age     3  0.00218115 0.016258  0.134157 8.9328e-01
##   treat_currentP     2 -0.01297930 0.020336 -0.638240 5.2332e-01
##   treat_currentP     3  0.00022289 0.021704  0.010269 9.9181e-01
##   treat_currentP     4 -0.03876031 0.020154 -1.923231 5.4451e-02
##   treat_currentP     5 -0.04926600 0.020614 -2.389920 1.6852e-02   *
##     treat_gender     2 -0.00518206 0.012479 -0.415263 6.7795e-01
##       treat_lawS     2 -0.01972457 0.018764 -1.051194 2.9317e-01
##       treat_lawS     3 -0.04386852 0.018859 -2.326152 2.0010e-02   *
##       treat_lawS     4 -0.08701392 0.018071 -4.815162 1.4708e-06 ***
##  treat_pAbortion     2 -0.09254834 0.017174 -5.388778 7.0938e-08 ***
##  treat_pAbortion     3 -0.08346417 0.018725 -4.457367 8.2973e-06 ***
##       treat_race     2  0.08244811 0.018797  4.386291 1.1530e-05 ***
##       treat_race     3  0.03215455 0.018638  1.725198 8.4492e-02
## ---
## Number of Obs. = 9947
## ---
## Number of Respondents = 2499
## ---
## ------------------------------------------------------------
## Conditional AMCE's (judicialknowledgehigh = 0):
## ------------------------------------------------------------
##    Attribute Level Estimate Std. Err z value Pr(>|z|)
##  same.gender     1 0.021789 0.015738  1.3845   0.1662
## ---
## Number of Obs. = 9947
## Number of Respondents = 2499
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05
##
## ------------------------------------------------------------
## Conditional AMCE's (judicialknowledgehigh = 1):
## ------------------------------------------------------------
##    Attribute Level   Estimate Std. Err  z value Pr(>|z|)
##  same.gender     1 -0.0069956 0.019478 -0.35915  0.71948
## ---
## Number of Obs. = 9947
## Number of Respondents = 2499
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05
##
```

```
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05
##
## --------------------
## AMCE Baseline Levels:
## --------------------
##        Attribute Level
##      politicized
##      same.gender
##        same.race
##        treat_age
##   treat_currentP
##     treat_gender
##       treat_lawS
##  treat_pAbortion
##       treat_race
```

```
results.support.knowledge.race <- amce(support2 ~ politicized + treat_age + treat_gender +
                                       treat_race + same.gender +
                                       same.race*judicial.knowledge.high + treat_lawS +
                                       treat_currentP + treat_pAbortion,
                                       data=data.support, cluster=TRUE,
                                       respondent.id="caseid", weights="weight",
                                       respondent.varying="judicial.knowledge.high")
summary(results.support.knowledge.race)
```

```
## ------------------------------------------
## Average Marginal Component Effects (AMCE):
## ------------------------------------------
##        Attribute Level    Estimate Std. Err   z value   Pr(>|z|)
##      politicized     1 -0.02152808 0.016788 -1.282321 1.9973e-01
##      same.gender     1  0.01267180 0.012395  1.022313 3.0663e-01
##        same.race     1  0.06145415 0.017882  3.436691 5.8887e-04 ***
##        treat_age     2  0.01262425 0.016684  0.756663 4.4925e-01
##        treat_age     3  0.00218115 0.016258  0.134157 8.9328e-01
##   treat_currentP     2 -0.01297930 0.020336 -0.638240 5.2332e-01
##   treat_currentP     3  0.00022289 0.021704  0.010269 9.9181e-01
##   treat_currentP     4 -0.03876031 0.020154 -1.923231 5.4451e-02
##   treat_currentP     5 -0.04926600 0.020614 -2.389920 1.6852e-02   *
##     treat_gender     2 -0.00518206 0.012479 -0.415263 6.7795e-01
##       treat_lawS     2 -0.01972457 0.018764 -1.051194 2.9317e-01
##       treat_lawS     3 -0.04386852 0.018859 -2.326152 2.0010e-02   *
##       treat_lawS     4 -0.08701392 0.018071 -4.815162 1.4708e-06 ***
##  treat_pAbortion     2 -0.09254834 0.017174 -5.388778 7.0938e-08 ***
##  treat_pAbortion     3 -0.08346417 0.018725 -4.457367 8.2973e-06 ***
##       treat_race     2  0.08244811 0.018797  4.386291 1.1530e-05 ***
##       treat_race     3  0.03215455 0.018638  1.725198 8.4492e-02
## ---
## Number of Obs. = 9947
## ---
## Number of Respondents = 2499
## ---
## ------------------------------------------------------------
## Conditional AMCE's (judicialknowledgehigh = 0):
## ------------------------------------------------------------
##  Attribute Level Estimate Std. Err z value  Pr(>|z|)
```

```
##  same.race     1 0.055914 0.019324  2.8935 0.0038093 **
## ---
## Number of Obs. = 9947
## Number of Respondents = 2499
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05
##
## ----------------------------------------------------------------
## Conditional AMCE's (judicialknowledgehigh = 1):
## ----------------------------------------------------------------
##  Attribute Level Estimate Std. Err z value  Pr(>|z|)
##  same.race     1 0.075219 0.027919  2.6941 0.0070571 **
## ---
## Number of Obs. = 9947
## Number of Respondents = 2499
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05
##
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05
##
## --------------------
## AMCE Baseline Levels:
## --------------------
##        Attribute Level
##      politicized
##      same.gender
##        same.race
##        treat_age
##   treat_currentP
##     treat_gender
##       treat_lawS
##  treat_pAbortion
##       treat_race
```

```
# Coefficients and SEs for support and knowledge interaction
se_gender_low <- 0.015738
se_race_low <- 0.019324
se_gender_high <-  0.019478
se_race_high <- 0.027919

coef_gender_low <- 0.021789
coef_race_low <- 0.055914
coef_gender_high <- -0.0069956
coef_race_high <-0.075219


# Create the confidence intervals
z_95 <- qnorm(.975)
ci_1 <- c(coef_gender_low  - z_95*se_gender_low,
          coef_gender_low  + z_95*se_gender_low)
ci_2 <- c(coef_race_low    - z_95*se_race_low,
          coef_race_low    + z_95*se_race_low)
ci_3 <- c(coef_gender_high - z_95*se_gender_high,
          coef_gender_high + z_95*se_gender_high)
```

```r
ci_4 <- c(coef_race_high   - z_95*se_race_high,
          coef_race_high   + z_95*se_race_high)

gender_low <- data.frame(est = coef_gender_low, lb = ci_1[1],
                         ub = ci_1[2], model = "Gender (Low Knowledge)")
race_low <- data.frame(est = coef_race_low, lb = ci_2[1],
                         ub = ci_2[2], model = "Race (Low Knowledge)")
gender_high <- data.frame(est = coef_gender_high, lb = ci_3[1],
                         ub = ci_3[2], model = "Gender (High Knowledge)")
race_high <- data.frame(est = coef_race_high, lb = ci_4[1],
                         ub = ci_4[2], model = "Race (High Knowledge)")

# Make the plot
ggplot(gender_low, aes(x = model, y = est)) +
  geom_point(colour="black") +
  geom_errorbar(aes(ymin = lb, ymax = ub), width = 0.2, colour="black") +
  geom_hline(yintercept = 0, lty = 2) +
  xlab("") +
  ylab("Coefficient Estimate") +
  geom_point(data= race_low, aes(x = model, y = est),colour="black") +
  geom_errorbar(data=race_low, aes(ymin = lb, ymax = ub), width = 0.2, colour="black") +
  geom_point(data= race_high, aes(x = model, y = est),colour="black") +
  geom_errorbar(data=race_high, aes(ymin = lb, ymax = ub), width = 0.2, colour="black") +
  geom_point(data= gender_high, aes(x = model, y = est),colour="black") +
  geom_errorbar(data=gender_high, aes(ymin = lb, ymax = ub), width = 0.2, colour="black")+
  theme_bw() + coord_flip()
```

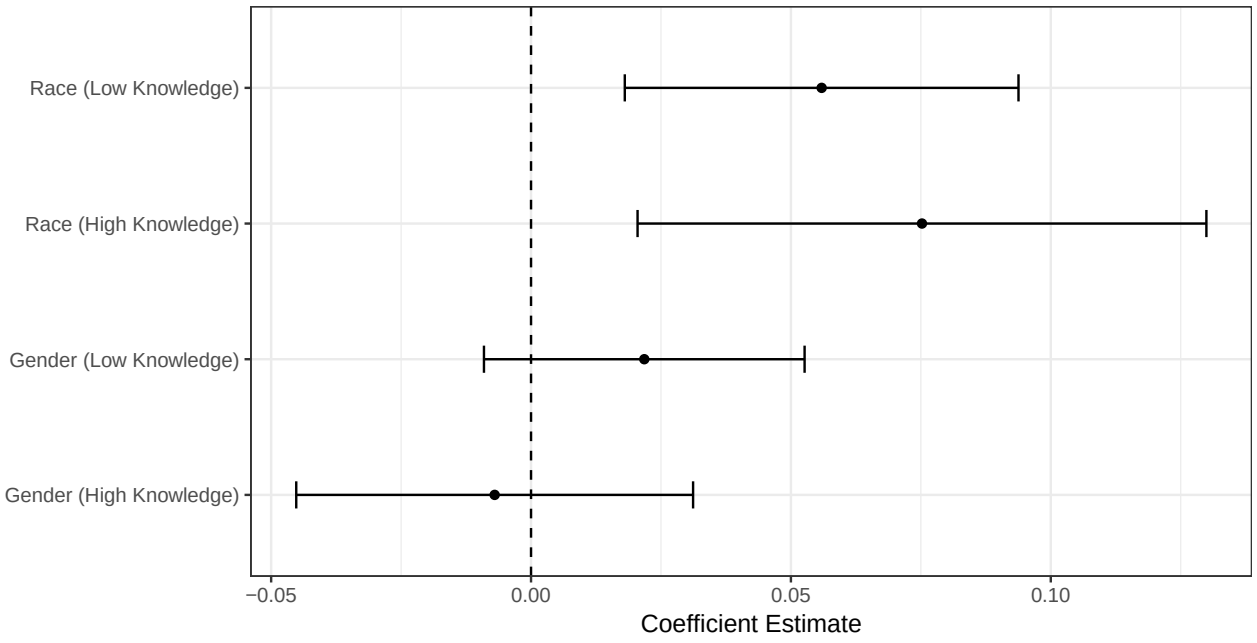


Figure A.3: Partisanship, Race and Trust in Nominee Impartiality (Binary Measure)

```
# Support estimates, white respondents
# Re-level the data to treat respondent's race/gender as the baseline for nominee evaluations
# Re Level the data to make sure that we are using respondent's race as the baseline
data.trust <- within(data.trust, race_nominee <- relevel(race_nominee,
                                                         ref = "White"))
data.trust <- within(data.trust, gender_nominee <- relevel(gender_nominee,
                                                         ref = "Male"))


# Trust, white respondents
data.trust.white <- data.trust[which(data.trust$race_respondent == "White"),]

# Democrats only
data.trust.white.dems <- data.trust.white[which(data.trust.white$pid3a == "Democrat"),]
results.trust.white.dems <- amce(trust2 ~ gender_nominee + race_nominee +
                                  politicized + treat_age + treat_lawS +
                                  treat_currentP + treat_pAbortion + same.gender,
                                  data=data.trust.white.dems,cluster=TRUE,
                                  respondent.id="caseid",weights="weight")

# Republicans only
data.trust.white.reps <- data.trust.white[which(data.trust.white$pid3a == "Republican"),]
results.trust.white.reps <- amce(trust2 ~ gender_nominee + race_nominee +
                                  politicized + treat_age +  treat_lawS +
                                  treat_currentP + treat_pAbortion + same.gender,
                                  data=data.trust.white.reps,cluster=TRUE,
                                  respondent.id="caseid",weights="weight")

# Trust, Black respondents

# Re Level the data to make sure that we are using respondent's race as the baseline
data.trust <- within(data.trust, race_nominee <- relevel(race_nominee,
                                                         ref = "Black"))
data.trust.black <- data.trust[which(data.trust$race_respondent == "Black"),]

# Democrats only
data.trust.black.dems <- data.trust.black[which(data.trust.black$pid3a == "Democrat"),]
results.trust.black.dems <- amce(trust2 ~ gender_nominee + race_nominee +
                                 politicized + treat_age + treat_lawS +
                                 treat_currentP + treat_pAbortion + same.gender,
                                 data=data.trust.black.dems,cluster=TRUE,
                                 respondent.id="caseid",weights="weight")

# Republicans only
data.trust.black.reps <- data.trust.black[which(data.trust.black$pid3a == "Republican"),]
results.trust.black.reps <- amce(trust2 ~ gender_nominee + race_nominee +
                                 politicized + treat_age + treat_lawS +
                                 treat_currentP + treat_pAbortion + same.gender,
                                 data=data.trust.black.reps,cluster=TRUE,
                                 respondent.id="caseid",weights="weight")
```

```r
# Trust, Hispanic respondents

# Re Level the data to make sure that we are using respondent's race as the baseline
data.trust <- within(data.trust, race_nominee <- relevel(race_nominee,
                                                         ref = "Hispanic"))
yougov.hispanic <- data[which(data$race_respondent == "Hispanic"),]

data.trust.hispanic <- data.trust[which(data.trust$race_respondent == "Hispanic"),]

# Democrats only
data.trust.hispanic.dems <- data.trust.hispanic[which(data.trust.hispanic$pid3a == "Democrat"),]
results.trust.hispanic.dems <- amce(trust2 ~ gender_nominee + race_nominee +
                                    politicized + treat_age + treat_lawS +
                                    treat_currentP + treat_pAbortion + same.gender,
                                    data=data.trust.hispanic.dems,cluster=TRUE,
                                    respondent.id="caseid",weights="weight")

# Republicans only
data.trust.hispanic.reps <- data.trust.hispanic[which(data.trust.hispanic$pid3a == "Republican"),]
results.trust.hispanic.reps <- amce(trust2 ~ gender_nominee + race_nominee +
                                    politicized + treat_age +  treat_lawS +
                                    treat_currentP + treat_pAbortion + same.gender,
                                    data=data.trust.hispanic.reps,cluster=TRUE,
                                    respondent.id="caseid",weights="weight")

# Coefficients and SEs for Democrats
se.hispanicresp.dem.white <- 0.089825
se.hispanicresp.dem.black <- 0.078777

se.blackresp.dem.hispanic <- 0.058512
se.blackresp.dem.white <- 0.060110

se.whiteresp.dem.hispanic <- 0.024663
se.whiteresp.dem.black <-  0.031842

coef.hispanicresp.dem.white <- 0.069204
coef.hispanicresp.dem.black <- -0.043731

coef.blackresp.dem.hispanic <- -0.1364005
coef.blackresp.dem.white <- -0.2345870

coef.whiteresp.dem.hispanic <- 0.0380279
coef.whiteresp.dem.black <- 0.0770066


# Create the confidence intervals
z_95 <- qnorm(.975)
ci_1 <- c(coef.hispanicresp.dem.white - z_95*se.hispanicresp.dem.white,
          coef.hispanicresp.dem.white + z_95*se.hispanicresp.dem.white)
ci_2 <- c(coef.hispanicresp.dem.black - z_95*se.hispanicresp.dem.black,
          coef.hispanicresp.dem.black + z_95*se.hispanicresp.dem.black)
ci_3 <- c(coef.blackresp.dem.hispanic - z_95*se.blackresp.dem.hispanic,
          coef.blackresp.dem.hispanic + z_95*se.blackresp.dem.hispanic)
```

```
ci_4 <- c(coef.blackresp.dem.white - z_95*se.blackresp.dem.white,
          coef.blackresp.dem.white + z_95*se.blackresp.dem.white)
ci_5 <- c(coef.whiteresp.dem.hispanic - z_95*se.whiteresp.dem.hispanic,
          coef.whiteresp.dem.hispanic + z_95*se.whiteresp.dem.hispanic)
ci_6 <- c(coef.whiteresp.dem.black - z_95*se.whiteresp.dem.black,
          coef.whiteresp.dem.black + z_95*se.whiteresp.dem.black)

hispanicresp.dem.white <- data.frame(est = coef.hispanicresp.dem.white,
                                     lb = ci_1[1], ub = ci_1[2], model = "Hispanic-White")
hispanicresp.dem.black <- data.frame(est = coef.hispanicresp.dem.black,
                                     lb = ci_2[1], ub = ci_2[2], model = "Hispanic-Black")
blackresp.dem.hispanic <- data.frame(est = coef.blackresp.dem.hispanic,
                                     lb = ci_3[1], ub = ci_3[2], model = "Black-Hispanic")
blackresp.dem.white <- data.frame(est = coef.blackresp.dem.white,
                                  lb = ci_4[1], ub = ci_4[2], model = "Black-White")
whiteresp.dem.hispanic <- data.frame(est = coef.whiteresp.dem.hispanic,
                                     lb = ci_5[1], ub = ci_5[2], model = "White-Hispanic")
whiteresp.dem.black <- data.frame(est = coef.whiteresp.dem.black,
                                  lb = ci_6[1], ub = ci_6[2],  model = "White-Black")


# Coefficients and SEs for Republicans
se.hispanicresp.rep.white <- 0.119063
se.hispanicresp.rep.black <- 0.110455

se.blackresp.rep.hispanic <- 0.16202
se.blackresp.rep.white <- 0.17796

se.whiteresp.rep.hispanic <- 0.030534
se.whiteresp.rep.black <-  0.031648

coef.hispanicresp.rep.white <- -0.1642361
coef.hispanicresp.rep.black <- -0.0875098

coef.blackresp.rep.hispanic <- -0.312697
coef.blackresp.rep.white <- -0.094893

coef.whiteresp.rep.hispanic <- -0.016244
coef.whiteresp.rep.black <-  -0.053209


# Create the confidence intervals
ci_1 <- c(coef.hispanicresp.rep.white - z_95*se.hispanicresp.rep.white,
          coef.hispanicresp.rep.white + z_95*se.hispanicresp.rep.white)
ci_2 <- c(coef.hispanicresp.rep.black - z_95*se.hispanicresp.rep.black,
          coef.hispanicresp.rep.black + z_95*se.hispanicresp.rep.black)
ci_3 <- c(coef.blackresp.rep.hispanic - z_95*se.blackresp.rep.hispanic,
          coef.blackresp.rep.hispanic + z_95*se.blackresp.rep.hispanic)
ci_4 <- c(coef.blackresp.rep.white - z_95*se.blackresp.rep.white,
          coef.blackresp.rep.white + z_95*se.blackresp.rep.white)
ci_5 <- c(coef.whiteresp.rep.hispanic - z_95*se.whiteresp.rep.hispanic,
          coef.whiteresp.rep.hispanic + z_95*se.whiteresp.rep.hispanic)
ci_6 <- c(coef.whiteresp.rep.black - z_95*se.whiteresp.rep.black,
```

```
          coef.whiteresp.rep.black + z_95*se.whiteresp.rep.black)

hispanicresp.rep.white <- data.frame(est = coef.hispanicresp.rep.white,
                                     lb = ci_1[1], ub = ci_1[2], model = "Hispanic-White")
hispanicresp.rep.black <- data.frame(est = coef.hispanicresp.rep.black,
                                     lb = ci_2[1], ub = ci_2[2], model = "Hispanic-Black")
blackresp.rep.hispanic <- data.frame(est = coef.blackresp.rep.hispanic,
                                     lb = ci_3[1], ub = ci_3[2], model = "Black-Hispanic")
blackresp.rep.white <- data.frame(est = coef.blackresp.rep.white,
                                     lb = ci_4[1], ub = ci_4[2], model = "Black-White")
whiteresp.rep.hispanic <- data.frame(est = coef.whiteresp.rep.hispanic,
                                     lb = ci_5[1], ub = ci_5[2], model = "White-Hispanic")
whiteresp.rep.black <- data.frame(est = coef.whiteresp.rep.black,
                                     lb = ci_6[1], ub = ci_6[2],  model = "White-Black")

# Democrats
g1 <- ggplot(hispanicresp.dem.white, aes(x = model, y = est)) +
  geom_point(colour="grey45") +
  geom_errorbar(aes(ymin = lb, ymax = ub), width = 0.2, colour="grey45") +
  geom_hline(yintercept = 0, lty = 2) +
  xlab("") +
  ylab("") +
  geom_point(data= hispanicresp.dem.black, aes(x = model, y = est),colour="grey45") +
  geom_errorbar(data=hispanicresp.dem.black, aes(ymin = lb, ymax = ub), width = 0.2,
  colour="grey45") +
  geom_point(data= blackresp.dem.hispanic, aes(x = model, y = est),colour="grey70") +
  geom_errorbar(data=blackresp.dem.hispanic, aes(ymin = lb, ymax = ub), width = 0.2,
  colour="grey70") +
  geom_point(data= blackresp.dem.white, aes(x = model, y = est),colour="grey70") +
  geom_errorbar(data=blackresp.dem.white, aes(ymin = lb, ymax = ub), width = 0.2,
  colour="grey70")  +
  geom_point(data= whiteresp.dem.hispanic, aes(x = model, y = est),colour="black") +
  geom_errorbar(data=whiteresp.dem.hispanic, aes(ymin = lb, ymax = ub), width = 0.2,
  colour="black") +
  geom_point(data= whiteresp.dem.black, aes(x = model, y = est),colour="black") +
  geom_errorbar(data=whiteresp.dem.black, aes(ymin = lb, ymax = ub), width = 0.2,
  colour="black")  +
  theme_bw() + coord_flip() + ylim(-.7,0.5) +
  labs(title="Democrats: Trust in Nominee Impartiality by Race") +
  theme(plot.title = element_text(hjust = 0.5))

# Republicans
g2 <- ggplot(hispanicresp.rep.white, aes(x = model, y = est)) +
  geom_point(colour="grey45") +
  geom_errorbar(aes(ymin = lb, ymax = ub), width = 0.2, colour="grey45") +
  geom_hline(yintercept = 0, lty = 2) +
  xlab("") +
  ylab("") +
  geom_point(data= hispanicresp.rep.black, aes(x = model, y = est),colour="grey45") +
  geom_errorbar(data=hispanicresp.rep.black, aes(ymin = lb, ymax = ub), width = 0.2,
  colour="grey45") +
  geom_point(data= blackresp.rep.hispanic, aes(x = model, y = est),colour="grey70") +
  geom_errorbar(data=blackresp.rep.hispanic, aes(ymin = lb, ymax = ub), width = 0.2,
```

```
  colour="grey70") +
  geom_point(data= blackresp.rep.white, aes(x = model, y = est),colour="grey70") +
  geom_errorbar(data=blackresp.rep.white, aes(ymin = lb, ymax = ub), width = 0.2,
  colour="grey70")  +
  geom_point(data= whiteresp.rep.hispanic, aes(x = model, y = est),colour="black") +
  geom_errorbar(data=whiteresp.rep.hispanic, aes(ymin = lb, ymax = ub), width = 0.2,
  colour="black") +
  geom_point(data= whiteresp.rep.black, aes(x = model, y = est),colour="black") +
  geom_errorbar(data=whiteresp.rep.black, aes(ymin = lb, ymax = ub), width = 0.2,
  colour="black")  +
  theme_bw() + coord_flip()  + ylim(-.7,0.5) +
  labs(title="Republicans: Trust in Nominee Impartiality by Race") +
  theme(plot.title = element_text(hjust = 0.5))

grid.arrange(g1, g2, ncol=1, left= textGrob("Respondent-Candidate Pairing", rot=90),
             bottom=textGrob("Coefficient Estimate"))
```

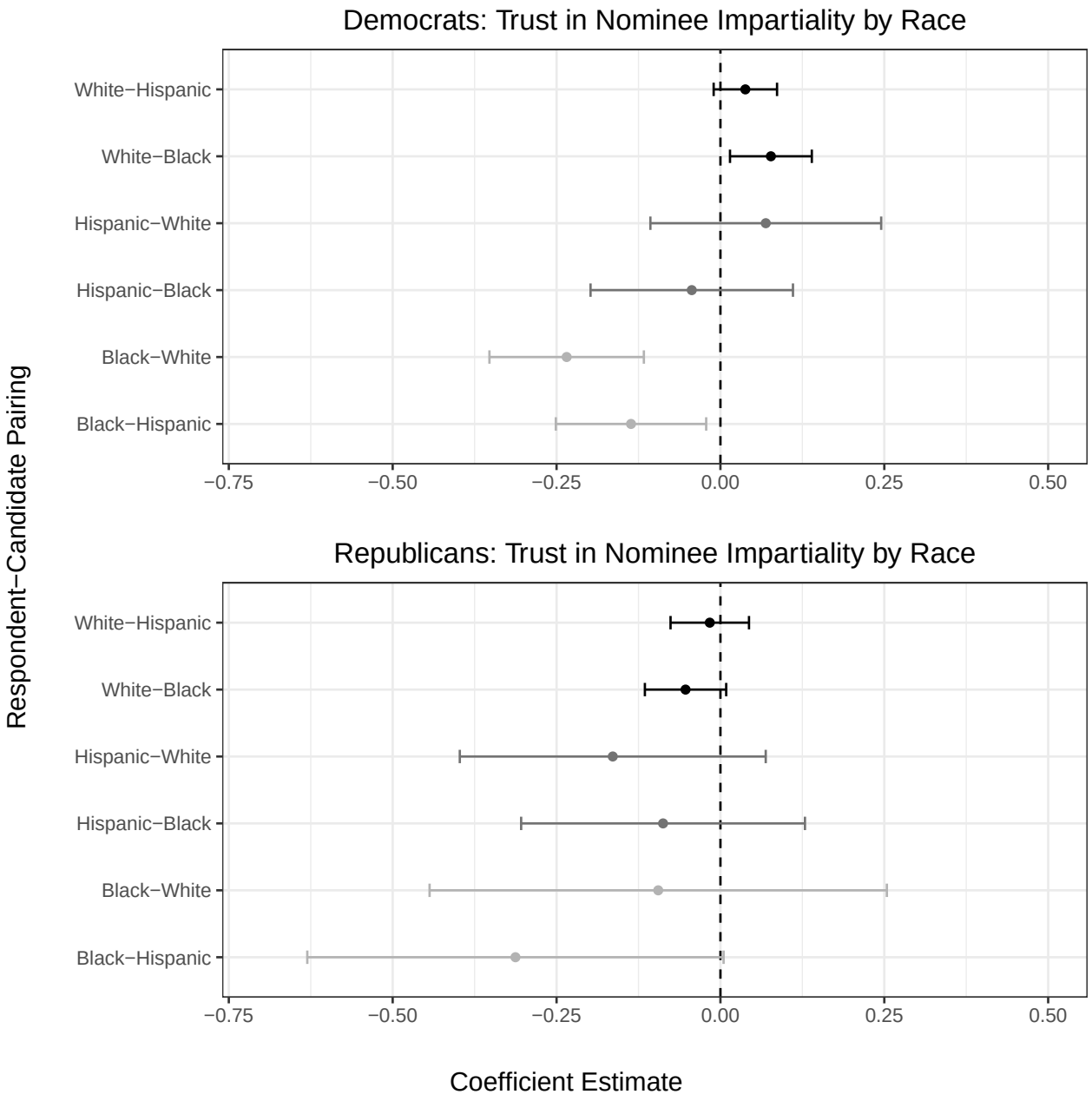


Figure A.4: Partisanship, Race and Nominee Qualifications (Binary Measure)

```
# Re Level the data to make sure that we are using respondent's race as the baseline
data.qualified <- within(data.qualified,
                         race_nominee <- relevel(race_nominee, ref = "White"))
data.qualified <- within(data.qualified,
                         gender_nominee <- relevel(gender_nominee, ref = "Male"))


# Qualified, white respondents
```

```
data.qualified.white <- data.qualified[which(data.qualified$race_respondent == "White"),]

# Democrats only
data.qualified.white.dems <- data.qualified.white[which(data.qualified.white$pid3a ==
                                                        "Democrat"),]
results.qualified.white.dems <- amce(qualified2 ~ gender_nominee + race_nominee +
                                     politicized + treat_age + treat_lawS + treat_currentP +
                                     treat_pAbortion + same.gender,
                                     data=data.qualified.white.dems,cluster=TRUE,
                                     respondent.id="caseid",weights="weight")

# Republicans only
data.qualified.white.reps <- data.qualified.white[which(data.qualified.white$pid3a ==
                                                        "Republican"),]
results.qualified.white.reps <- amce(qualified2 ~ gender_nominee + race_nominee +
                                     politicized + treat_age +  treat_lawS +
                                     treat_currentP + treat_pAbortion + same.gender,
                                     data=data.qualified.white.reps,cluster=TRUE,
                                     respondent.id="caseid",weights="weight")

# Qualified, Black respondents

# Re Level the data to make sure that we are using respondent's race as the baseline
data.qualified <- within(data.qualified, race_nominee <- relevel(race_nominee,
                                                                 ref = "Black"))
data.qualified.black <- data.qualified[which(data.qualified$race_respondent == "Black"),]


# Democrats only
data.qualified.black.dems <- data.qualified.black[which(data.qualified.black$pid3a ==
                                                        "Democrat"),]
results.qualified.black.dems <- amce(qualified2 ~ gender_nominee + race_nominee +
                                     politicized + treat_age + treat_lawS +
                                     treat_currentP + treat_pAbortion + same.gender,
                                     data=data.qualified.black.dems,cluster=TRUE,
                                     respondent.id="caseid",weights="weight")

# Republicans only
data.qualified.black.reps <- data.qualified.black[which(data.qualified.black$pid3a ==
                                                        "Republican"),]
results.qualified.black.reps <- amce(qualified2 ~ gender_nominee + race_nominee +
                                     politicized + treat_age + treat_lawS +
                                     treat_currentP + treat_pAbortion + same.gender,
                                     data=data.qualified.black.reps,cluster=TRUE,
                                     respondent.id="caseid",weights="weight")

# Qualified, Hispanic respondents

# Re Level the data to make sure that we are using respondent's race as the baseline
data.qualified <- within(data.qualified, race_nominee <- relevel(race_nominee,
                                                                 ref = "Hispanic"))
data.qualified.hispanic <- data.qualified[which(data.qualified$race_respondent ==
                                                  "Hispanic"),]
```

```
# Democrats only
data.qualified.hispanic.dems <- data.qualified.hispanic[which(data.qualified.hispanic$pid3a ==
                                                              "Democrat"),]
results.qualified.hispanic.dems <- amce(qualified2 ~ gender_nominee + race_nominee +
                                        politicized + treat_age + treat_lawS +
                                        treat_currentP + treat_pAbortion + same.gender,
                                        data=data.qualified.hispanic.dems,cluster=TRUE,
                                        respondent.id="caseid",weights="weight")

# Republicans only
data.qualified.hispanic.reps <- data.qualified.hispanic[which(data.qualified.hispanic$pid3a ==
                                                              "Republican"),]
results.qualified.hispanic.reps <- amce(qualified2 ~ gender_nominee + race_nominee +
                                        politicized + treat_age +  treat_lawS +
                                        treat_currentP + treat_pAbortion + same.gender,
                                        data=data.qualified.hispanic.reps,cluster=TRUE,
                                        respondent.id="caseid",weights="weight")

# Coefficients and SEs for Democrats
se.hispanicresp.dem.white <- 0.065766
se.hispanicresp.dem.black <- 0.077770

se.blackresp.dem.hispanic <- 0.072084
se.blackresp.dem.white <- 0.050053

se.whiteresp.dem.hispanic <- 0.031023
se.whiteresp.dem.black <-  0.036913

coef.hispanicresp.dem.white <- -0.0687146
coef.hispanicresp.dem.black <- -0.0066937

coef.blackresp.dem.hispanic <- -0.07991862
coef.blackresp.dem.white <- -0.15624990

coef.whiteresp.dem.hispanic <- 0.0249701
coef.whiteresp.dem.black <- 0.0496341


# Create the confidence intervals
ci_1 <- c(coef.hispanicresp.dem.white - z_95*se.hispanicresp.dem.white,
          coef.hispanicresp.dem.white + z_95*se.hispanicresp.dem.white)
ci_2 <- c(coef.hispanicresp.dem.black - z_95*se.hispanicresp.dem.black,
          coef.hispanicresp.dem.black + z_95*se.hispanicresp.dem.black)
ci_3 <- c(coef.blackresp.dem.hispanic - z_95*se.blackresp.dem.hispanic,
          coef.blackresp.dem.hispanic + z_95*se.blackresp.dem.hispanic)
ci_4 <- c(coef.blackresp.dem.white    - z_95*se.blackresp.dem.white,
          coef.blackresp.dem.white    + z_95*se.blackresp.dem.white)
ci_5 <- c(coef.whiteresp.dem.hispanic - z_95*se.whiteresp.dem.hispanic,
          coef.whiteresp.dem.hispanic + z_95*se.whiteresp.dem.hispanic)
ci_6 <- c(coef.whiteresp.dem.black    - z_95*se.whiteresp.dem.black,
          coef.whiteresp.dem.black    + z_95*se.whiteresp.dem.black)

hispanicresp.dem.white <- data.frame(est = coef.hispanicresp.dem.white,
```

```
                                lb = ci_1[1], ub = ci_1[2], model = "Hispanic-White")
hispanicresp.dem.black <- data.frame(est = coef.hispanicresp.dem.black,
                                lb = ci_2[1], ub = ci_2[2], model = "Hispanic-Black")
blackresp.dem.hispanic <- data.frame(est = coef.blackresp.dem.hispanic,
                                lb = ci_3[1], ub = ci_3[2], model = "Black-Hispanic")
blackresp.dem.white <- data.frame(est = coef.blackresp.dem.white,
                                lb = ci_4[1], ub = ci_4[2], model = "Black-White")
whiteresp.dem.hispanic <- data.frame(est = coef.whiteresp.dem.hispanic,
                                lb = ci_5[1], ub = ci_5[2], model = "White-Hispanic")
whiteresp.dem.black <- data.frame(est = coef.whiteresp.dem.black,
                                lb = ci_6[1], ub = ci_6[2], model = "White-Black")


# Coefficients and SEs for Republicans
se.hispanicresp.rep.white <- 0.114892
se.hispanicresp.rep.black <- 0.111226

se.blackresp.rep.hispanic <- 0.137968
se.blackresp.rep.white <- 0.122147

se.whiteresp.rep.hispanic <- 0.033191
se.whiteresp.rep.black <-  0.031223

coef.hispanicresp.rep.white <- -0.2200228
coef.hispanicresp.rep.black <- -0.0017158

coef.blackresp.rep.hispanic <- -0.527848
coef.blackresp.rep.white <- -0.346457

coef.whiteresp.rep.hispanic <- -0.0187308
coef.whiteresp.rep.black <- -0.0768528


# Create the confidence intervals
ci_1 <- c(coef.hispanicresp.rep.white - z_95*se.hispanicresp.rep.white,
          coef.hispanicresp.rep.white + z_95*se.hispanicresp.rep.white)
ci_2 <- c(coef.hispanicresp.rep.black - z_95*se.hispanicresp.rep.black,
          coef.hispanicresp.rep.black + z_95*se.hispanicresp.rep.black)
ci_3 <- c(coef.blackresp.rep.hispanic - z_95*se.blackresp.rep.hispanic,
          coef.blackresp.rep.hispanic + z_95*se.blackresp.rep.hispanic)
ci_4 <- c(coef.blackresp.rep.white    - z_95*se.blackresp.rep.white,
          coef.blackresp.rep.white    + z_95*se.blackresp.rep.white)
ci_5 <- c(coef.whiteresp.rep.hispanic - z_95*se.whiteresp.rep.hispanic,
          coef.whiteresp.rep.hispanic + z_95*se.whiteresp.rep.hispanic)
ci_6 <- c(coef.whiteresp.rep.black    - z_95*se.whiteresp.rep.black,
          coef.whiteresp.rep.black    + z_95*se.whiteresp.rep.black)

hispanicresp.rep.white <- data.frame(est = coef.hispanicresp.rep.white,
                                lb = ci_1[1], ub = ci_1[2], model = "Hispanic-White")
hispanicresp.rep.black <- data.frame(est = coef.hispanicresp.rep.black,
                                lb = ci_2[1], ub = ci_2[2], model = "Hispanic-Black")
blackresp.rep.hispanic <- data.frame(est = coef.blackresp.rep.hispanic,
                                lb = ci_3[1], ub = ci_3[2], model = "Black-Hispanic")
```

```
blackresp.rep.white <- data.frame(est = coef.blackresp.rep.white,
                                  lb = ci_4[1], ub = ci_4[2],  model = "Black-White")
whiteresp.rep.hispanic <- data.frame(est = coef.whiteresp.rep.hispanic,
                                  lb = ci_5[1], ub = ci_5[2], model = "White-Hispanic")
whiteresp.rep.black <- data.frame(est = coef.whiteresp.rep.black,
                                  lb = ci_6[1], ub = ci_6[2], model = "White-Black")

# Make the plot

# Democrats
g1 <- ggplot(hispanicresp.dem.white, aes(x = model, y = est)) +
  geom_point(colour="grey45") +
  geom_errorbar(aes(ymin = lb, ymax = ub), width = 0.2, colour="grey45") +
  geom_hline(yintercept = 0, lty = 2) +
  xlab("") +
  ylab("") +
  geom_point(data= hispanicresp.dem.black, aes(x = model, y = est),colour="grey45") +
  geom_errorbar(data=hispanicresp.dem.black, aes(ymin = lb, ymax = ub), width = 0.2,
  colour="grey45") +
  geom_point(data= blackresp.dem.hispanic, aes(x = model, y = est),colour="grey70") +
  geom_errorbar(data=blackresp.dem.hispanic, aes(ymin = lb, ymax = ub), width = 0.2,
  colour="grey70") +
  geom_point(data= blackresp.dem.white, aes(x = model, y = est),colour="grey70") +
  geom_errorbar(data=blackresp.dem.white, aes(ymin = lb, ymax = ub), width = 0.2,
  colour="grey70")  +
  geom_point(data= whiteresp.dem.hispanic, aes(x = model, y = est),colour="black") +
  geom_errorbar(data=whiteresp.dem.hispanic, aes(ymin = lb, ymax = ub), width = 0.2,
  colour="black") +
  geom_point(data= whiteresp.dem.black, aes(x = model, y = est),colour="black") +
  geom_errorbar(data=whiteresp.dem.black, aes(ymin = lb, ymax = ub), width = 0.2,
  colour="black")  +
  theme_bw() + coord_flip() + ylim(-.9,0.5) +
  labs(title="Democrats: Nominee Qualifications by Race") +
  theme(plot.title = element_text(hjust = 0.5))

# Republicans
g2 <- ggplot(hispanicresp.rep.white, aes(x = model, y = est)) +
  geom_point(colour="grey45") +
  geom_errorbar(aes(ymin = lb, ymax = ub), width = 0.2, colour="grey45") +
  geom_hline(yintercept = 0, lty = 2) +
  xlab("") +
  ylab("") +
  geom_point(data= hispanicresp.rep.black, aes(x = model, y = est),colour="grey45") +
  geom_errorbar(data=hispanicresp.rep.black, aes(ymin = lb, ymax = ub), width = 0.2,
  colour="grey45") +
  geom_point(data= blackresp.rep.hispanic, aes(x = model, y = est),colour="grey70") +
  geom_errorbar(data=blackresp.rep.hispanic, aes(ymin = lb, ymax = ub), width = 0.2,
  colour="grey70") +
  geom_point(data= blackresp.rep.white, aes(x = model, y = est),colour="grey70") +
  geom_errorbar(data=blackresp.rep.white, aes(ymin = lb, ymax = ub), width = 0.2,
  colour="grey70")  +
  geom_point(data= whiteresp.rep.hispanic, aes(x = model, y = est),colour="black") +
  geom_errorbar(data=whiteresp.rep.hispanic, aes(ymin = lb, ymax = ub), width = 0.2,
```

```
  colour="black") +
  geom_point(data= whiteresp.rep.black, aes(x = model, y = est),colour="black") +
  geom_errorbar(data=whiteresp.rep.black, aes(ymin = lb, ymax = ub), width = 0.2,
  colour="black")  +
  theme_bw() + coord_flip()  + ylim(-.9,0.5) +
  labs(title="Republicans: Nominee Qualifications by Race") +
  theme(plot.title = element_text(hjust = 0.5))

grid.arrange(g1, g2, ncol=1, left= textGrob("Respondent-Candidate Pairing", rot=90),
             bottom=textGrob("Coefficient Estimate"))
```

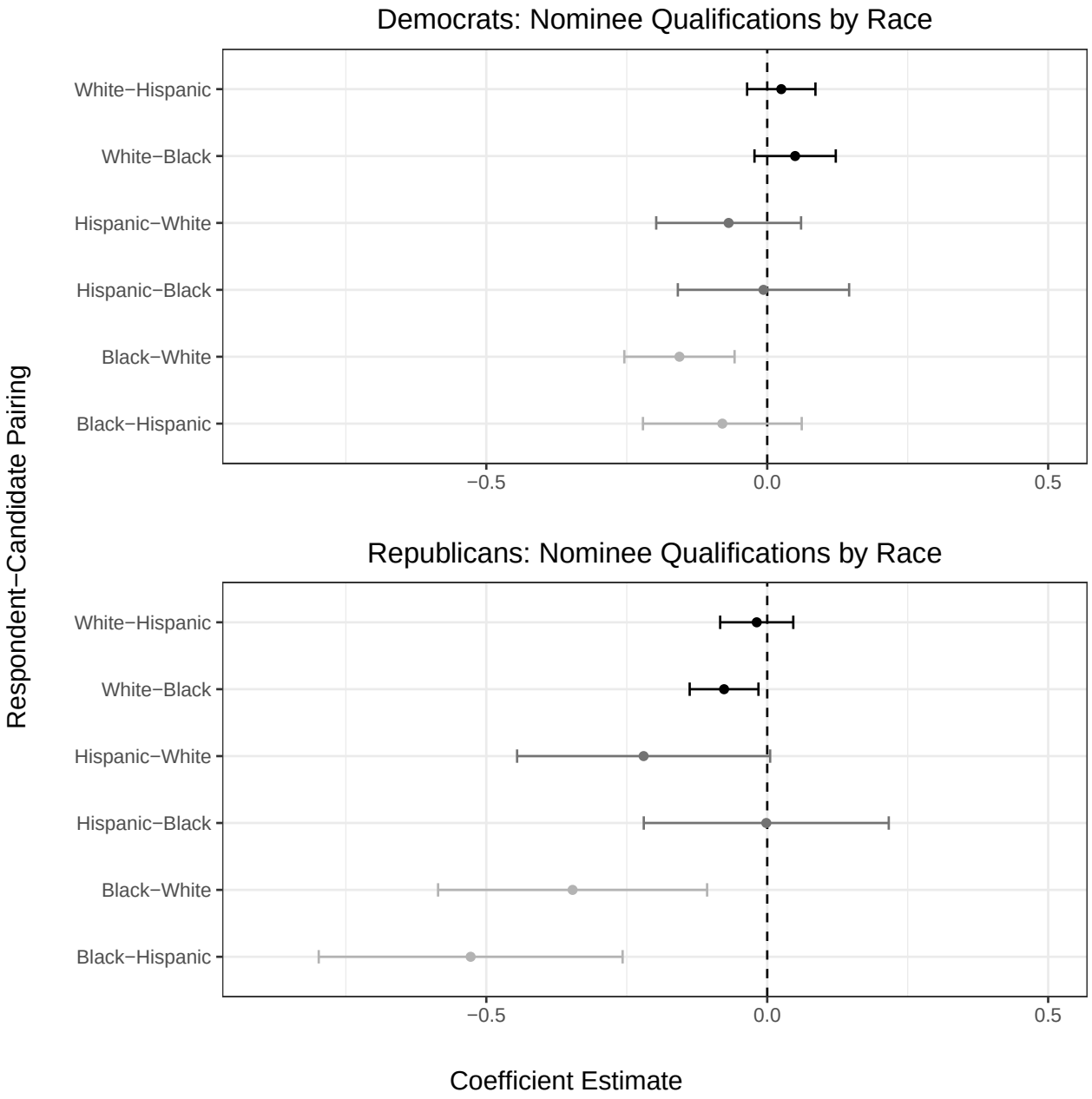


Figure A.5: Partisanship, Race, and Nominee Support

```
# Support, white respondents

# Re Level the data to make sure that we are using respondent's race as the baseline
data.support <- within(data.support, race_nominee <- relevel(race_nominee, ref = "White"))
data.support.white <- data.support[which(data.support$race_respondent == "White"),]

# Democrats only
data.support.white.dems <- data.support.white[which(data.support.white$pid3a == "Democrat"),]
results.support.white.dems <- amce(support2 ~ gender_nominee + race_nominee +
                                   politicized + treat_age + treat_lawS +
                                   treat_currentP + treat_pAbortion + same.gender,
                                   data=data.support.white.dems,cluster=TRUE,
                                   respondent.id="caseid",weights="weight")

# Republicans and Independents only
data.support.white.reps <- data.support.white[which(data.support.white$pid3a ==
                                       "Republican" | data.support.white$pid3a=="Independent"),]
results.support.white.reps <- amce(support2 ~ gender_nominee + race_nominee +
                                   politicized + treat_age +  treat_lawS +
                                   treat_currentP + treat_pAbortion + same.gender,
                                   data=data.support.white.reps,cluster=TRUE,
                                   respondent.id="caseid",weights="weight")

# Support, Black respondents

# Re Level the data to make sure that we are using respondent's race as the baseline
data.support <- within(data.support, race_nominee <- relevel(race_nominee, ref = "Black"))
data.support.black <- data.support[which(data.support$race_respondent == "Black"),]

# Democrats only
data.support.black.dems <- data.support.black[which(data.support.black$pid3a == "Democrat"),]
results.support.black.dems <- amce(support2 ~ gender_nominee + race_nominee +
                                   politicized + treat_age + treat_lawS +
                                   treat_currentP + treat_pAbortion + same.gender,
                                   data=data.support.black.dems,cluster=TRUE,
                                   respondent.id="caseid",weights="weight")

# Republicans and Independents only
data.support.black.reps <- data.support.black[which(data.support.black$pid3a ==
                                  "Republican" | data.support.black$pid3a=="Independent"),]
results.support.black.reps <- amce(support2 ~ gender_nominee + race_nominee +
                                   politicized + treat_age + treat_lawS +
                                   treat_currentP + treat_pAbortion + same.gender,
                                   data=data.support.black.reps,cluster=TRUE,
                                   respondent.id="caseid",weights="weight")

# Support, Hispanic respondents

# Re Level the data to make sure that we are using respondent's race as the baseline
data.support <- within(data.support, race_nominee <- relevel(race_nominee, ref = "Hispanic"))
data.support.hispanic <- data.support[which(data.support$race_respondent == "Hispanic"),]
```

```r
# Democrats only
data.support.hispanic.dems <- data.support.hispanic[which(data.support.hispanic$pid3a == "Democrat"),]
results.support.hispanic.dems <- amce(support2 ~ gender_nominee + race_nominee +
                                        politicized + treat_age + treat_lawS +
                                        treat_currentP + treat_pAbortion + same.gender,
                                        data=data.support.hispanic.dems,cluster=TRUE,
                                        respondent.id="caseid",weights="weight")

# Republicans and Independents only
data.support.hispanic.reps <- data.support.hispanic[which(data.support.hispanic$pid3a ==
                                      "Republican" |data.support.hispanic$pid3a=="Independent"),]
results.support.hispanic.reps <- amce(support2 ~ gender_nominee + race_nominee +
                                        politicized + treat_age +  treat_lawS +
                                        treat_currentP + treat_pAbortion + same.gender,
                                        data=data.support.hispanic.reps,cluster=TRUE,
                                        respondent.id="caseid",weights="weight")

# Coefficients and SEs for Democrats
se.hispanicresp.dem.white <- 0.084162
se.hispanicresp.dem.black <- 0.078847

se.blackresp.dem.hispanic <- 0.056423
se.blackresp.dem.white <- 0.054109

se.whiteresp.dem.hispanic <- 0.027374
se.whiteresp.dem.black <-  0.033561

coef.hispanicresp.dem.white <- -0.035620
coef.hispanicresp.dem.black <- 0.062250

coef.blackresp.dem.hispanic <- -0.1676413
coef.blackresp.dem.white <- -0.2253979

coef.whiteresp.dem.hispanic <- 0.0240293
coef.whiteresp.dem.black <- 0.0452267


# Create the confidence intervals
ci_1 <- c(coef.hispanicresp.dem.white - z_95*se.hispanicresp.dem.white,
          coef.hispanicresp.dem.white + z_95*se.hispanicresp.dem.white)
ci_2 <- c(coef.hispanicresp.dem.black - z_95*se.hispanicresp.dem.black,
          coef.hispanicresp.dem.black + z_95*se.hispanicresp.dem.black)
ci_3 <- c(coef.blackresp.dem.hispanic - z_95*se.blackresp.dem.hispanic,
          coef.blackresp.dem.hispanic + z_95*se.blackresp.dem.hispanic)
ci_4 <- c(coef.blackresp.dem.white    - z_95*se.blackresp.dem.white,
          coef.blackresp.dem.white    + z_95*se.blackresp.dem.white)
ci_5 <- c(coef.whiteresp.dem.hispanic - z_95*se.whiteresp.dem.hispanic,
          coef.whiteresp.dem.hispanic + z_95*se.whiteresp.dem.hispanic)
ci_6 <- c(coef.whiteresp.dem.black    - z_95*se.whiteresp.dem.black,
          coef.whiteresp.dem.black    + z_95*se.whiteresp.dem.black)

#graph the coefficients and their confidence intervals
hispanicresp.dem.white <- data.frame(est = coef.hispanicresp.dem.white,
```

```
                                  lb = ci_1[1], ub = ci_1[2], model = "Hispanic-White")
hispanicresp.dem.black <- data.frame(est = coef.hispanicresp.dem.black,
                                  lb = ci_2[1], ub = ci_2[2], model = "Hispanic-Black")
blackresp.dem.hispanic <- data.frame(est = coef.blackresp.dem.hispanic,
                                  lb = ci_3[1], ub = ci_3[2], model = "Black-Hispanic")
blackresp.dem.white <- data.frame(est = coef.blackresp.dem.white,
                                  lb = ci_4[1], ub = ci_4[2], model = "Black-White")
whiteresp.dem.hispanic <- data.frame(est = coef.whiteresp.dem.hispanic,
                                  lb = ci_5[1], ub = ci_5[2], model = "White-Hispanic")
whiteresp.dem.black <- data.frame(est = coef.whiteresp.dem.black,
                                  lb = ci_6[1], ub = ci_6[2], model = "White-Black")


# Coefficients and SEs for Republicans
se.hispanicresp.rep.white <- 0.049327
se.hispanicresp.rep.black <- 0.063217

se.blackresp.rep.hispanic <- 0.064957
se.blackresp.rep.white <- 0.073964

se.whiteresp.rep.hispanic <- 0.021805
se.whiteresp.rep.black <-  0.020299

coef.hispanicresp.rep.white <- -0.0252824
coef.hispanicresp.rep.black <-  0.1061138

coef.blackresp.rep.hispanic <- -0.0973707
coef.blackresp.rep.white <- -0.1136076

coef.whiteresp.rep.hispanic <- -0.0278392
coef.whiteresp.rep.black <-  -0.0095213


# Create the confidence intervals
ci_1 <- c(coef.hispanicresp.rep.white - z_95*se.hispanicresp.rep.white,
          coef.hispanicresp.rep.white + z_95*se.hispanicresp.rep.white)
ci_2 <- c(coef.hispanicresp.rep.black - z_95*se.hispanicresp.rep.black,
          coef.hispanicresp.rep.black + z_95*se.hispanicresp.rep.black)
ci_3 <- c(coef.blackresp.rep.hispanic - z_95*se.blackresp.rep.hispanic,
          coef.blackresp.rep.hispanic + z_95*se.blackresp.rep.hispanic)
ci_4 <- c(coef.blackresp.rep.white    - z_95*se.blackresp.rep.white,
          coef.blackresp.rep.white    + z_95*se.blackresp.rep.white)
ci_5 <- c(coef.whiteresp.rep.hispanic - z_95*se.whiteresp.rep.hispanic,
          coef.whiteresp.rep.hispanic + z_95*se.whiteresp.rep.hispanic)
ci_6 <- c(coef.whiteresp.rep.black    - z_95*se.whiteresp.rep.black,
          coef.whiteresp.rep.black    + z_95*se.whiteresp.rep.black)

hispanicresp.rep.white <- data.frame(est = coef.hispanicresp.rep.white,
                                  lb = ci_1[1], ub = ci_1[2], model = "Hispanic-White")
hispanicresp.rep.black <- data.frame(est = coef.hispanicresp.rep.black,
                                  lb = ci_2[1], ub = ci_2[2], model = "Hispanic-Black")
blackresp.rep.hispanic <- data.frame(est = coef.blackresp.rep.hispanic,
                                  lb = ci_3[1], ub = ci_3[2], model = "Black-Hispanic")
```

```r
blackresp.rep.white <- data.frame(est = coef.blackresp.rep.white,
                                  lb = ci_4[1], ub = ci_4[2], model = "Black-White")
whiteresp.rep.hispanic <- data.frame(est = coef.whiteresp.rep.hispanic,
                                  lb = ci_5[1], ub = ci_5[2], model = "White-Hispanic")
whiteresp.rep.black <- data.frame(est = coef.whiteresp.rep.black,
                                  lb = ci_6[1], ub = ci_6[2], model = "White-Black")

# Make the plot

# Democrats
g1 <- ggplot(hispanicresp.dem.white, aes(x = model, y = est)) +
  geom_point(colour="grey45") +
  geom_errorbar(aes(ymin = lb, ymax = ub), width = 0.2, colour="grey45") +
  geom_hline(yintercept = 0, lty = 2) +
  xlab("") +
  ylab("") +
  geom_point(data= hispanicresp.dem.black, aes(x = model, y = est),colour="grey45") +
  geom_errorbar(data=hispanicresp.dem.black, aes(ymin = lb, ymax = ub), width = 0.2,
  colour="grey45") +
  geom_point(data= blackresp.dem.hispanic, aes(x = model, y = est),colour="grey70") +
  geom_errorbar(data=blackresp.dem.hispanic, aes(ymin = lb, ymax = ub), width = 0.2,
  colour="grey70") +
  geom_point(data= blackresp.dem.white, aes(x = model, y = est),colour="grey70") +
  geom_errorbar(data=blackresp.dem.white, aes(ymin = lb, ymax = ub), width = 0.2,
  colour="grey70")  +
  geom_point(data= whiteresp.dem.hispanic, aes(x = model, y = est),colour="black") +
  geom_errorbar(data=whiteresp.dem.hispanic, aes(ymin = lb, ymax = ub), width = 0.2,
  colour="black") +
  geom_point(data= whiteresp.dem.black, aes(x = model, y = est),colour="black") +
  geom_errorbar(data=whiteresp.dem.black, aes(ymin = lb, ymax = ub), width = 0.2,
  colour="black")  +
  theme_bw() + coord_flip() + ylim(-.7,0.5) +
  labs(title="Democrats: Support for Nominees by Race") +
  theme(plot.title = element_text(hjust = 0.5))


# Republicans and Independents
g2 <- ggplot(hispanicresp.rep.white, aes(x = model, y = est)) +
  geom_point(colour="grey45") +
  geom_errorbar(aes(ymin = lb, ymax = ub), width = 0.2, colour="grey45") +
  geom_hline(yintercept = 0, lty = 2) +
  xlab("") +
  ylab("") +
  geom_point(data= hispanicresp.rep.black, aes(x = model, y = est),colour="grey45") +
  geom_errorbar(data=hispanicresp.rep.black, aes(ymin = lb, ymax = ub), width = 0.2,
  colour="grey45") +
  geom_point(data= blackresp.rep.hispanic, aes(x = model, y = est),colour="grey70") +
  geom_errorbar(data=blackresp.rep.hispanic, aes(ymin = lb, ymax = ub), width = 0.2,
  colour="grey70") +
  geom_point(data= blackresp.rep.white, aes(x = model, y = est),colour="grey70") +
  geom_errorbar(data=blackresp.rep.white, aes(ymin = lb, ymax = ub), width = 0.2,
  colour="grey70")  +
  geom_point(data= whiteresp.rep.hispanic, aes(x = model, y = est),colour="black") +
```

```
  geom_errorbar(data=whiteresp.rep.hispanic, aes(ymin = lb, ymax = ub), width = 0.2,
  colour="black") +
  geom_point(data= whiteresp.rep.black, aes(x = model, y = est),colour="black") +
  geom_errorbar(data=whiteresp.rep.black, aes(ymin = lb, ymax = ub), width = 0.2,
  colour="black")  +
  theme_bw() + coord_flip()  + ylim(-.7,0.5) +
  labs(title="Republicans and Independents: Support for Nominees by Race") +
  theme(plot.title = element_text(hjust = 0.5))

grid.arrange(g1, g2, ncol=1, left= textGrob("Respondent-Candidate Pairing", rot=90),
             bottom=textGrob("Coefficient Estimate"))
```

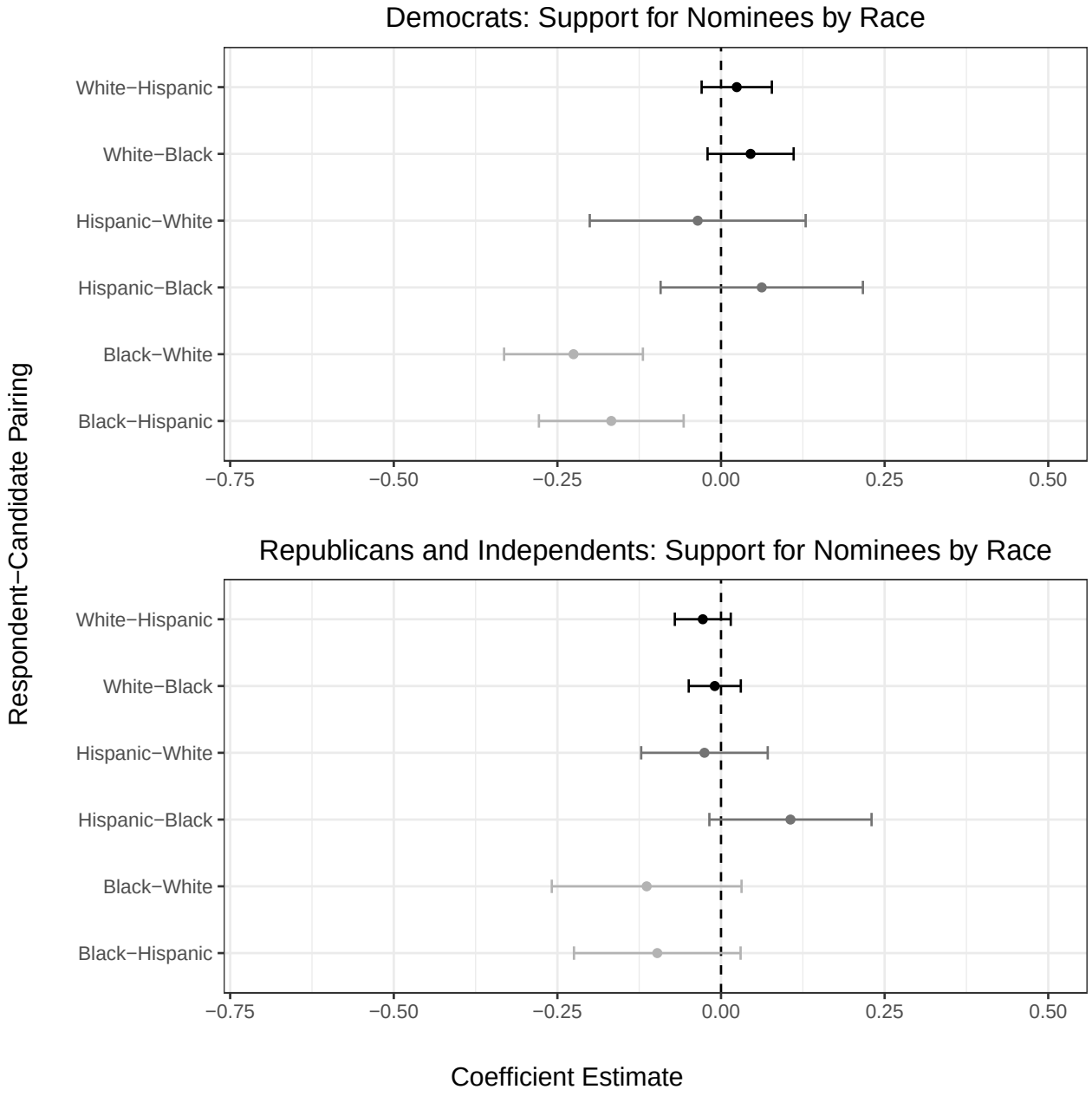


Figure A.6: Confidence in Supreme Court and Number of Female Justices

```
load("longitudinal_data.RData")

plot(confidence_women ~ year, data=confidence, pch=16, type="l", ylim=c(0.2,0.5), col="darkgreen",
     ylab="Percent Confident in Supreme Court", xlab="Year", xaxt="n", lwd=2, lty=0)
axis(side=1, at=c(1975,1980,1985,1990,1995,2000,2005,2010,2015), labels=c(1975,1980,1985,1990,1995,
                                                                          2000,2005,2010,2015))
polygon(x=c(1980,1980,1992,1992,1980), y=c(0.2,0.5,0.5,0.2,0.2), col="grey90")
polygon(x=c(1992,1992,2004,2004,1992), y=c(0.2,0.5,0.5,0.2,0.2), col="grey70")
polygon(x=c(2004,2004,2007,2007,2004), y=c(0.2,0.5,0.5,0.2,0.2), col="grey90")
polygon(x=c(2007,2007,2011,2011,2007), y=c(0.2,0.5,0.5,0.2,0.2), col="grey70")
polygon(x=c(2011,2011,2015,2015,2011), y=c(0.2,0.5,0.5,0.2,0.2), col="grey50")

lines(x=confidence$year, y=confidence$confidence_women, col="black", lwd=2, lty=2)
lines(x=confidence$year, y=confidence$confidence_men, col="darkblue", lwd=2)
text(x=1975, y=0.30, "Women",col="black", cex=1)
text(x=1975, y=0.42, "Men",col="darkblue", cex=1)

mtext(side=3, at=1980, text="O'Connor", cex=.75)
mtext(side=3, at=1992, text="Ginsburg", cex=.75)
mtext(side=3, at=2002, text="O'Connor Retires", cex=.75)
mtext(side=3, at=2007, text="Sotomayor", cex=.75)
mtext(side=3, at=2011, text="Kagan", cex=.75)
```

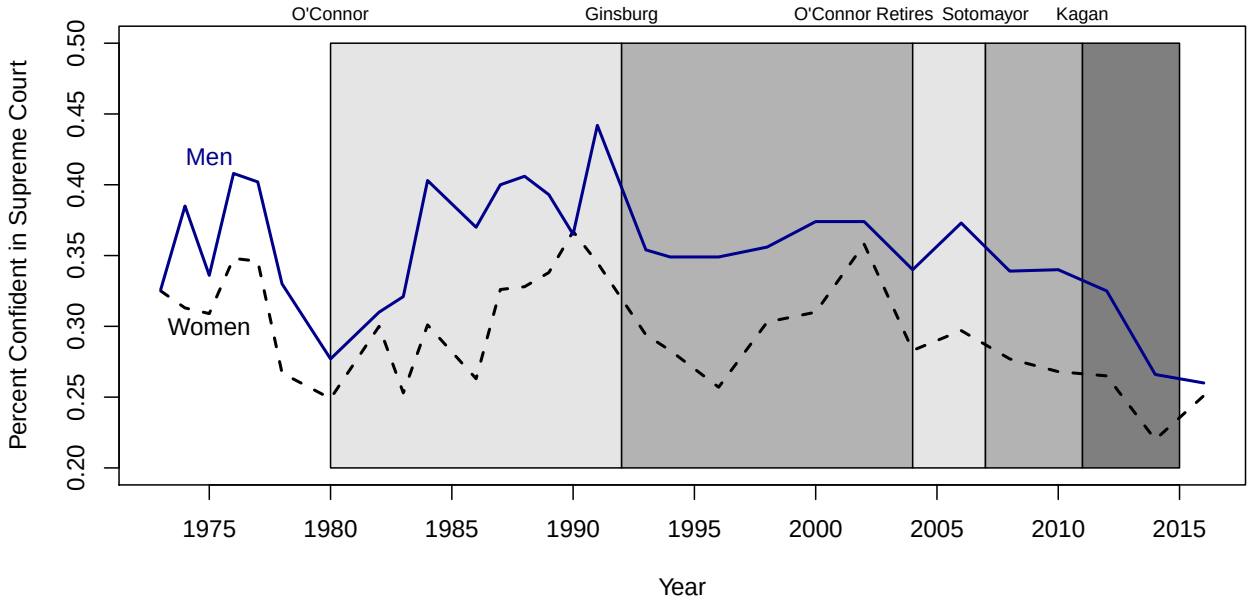


Table A.1: Interactive Relationship between Nominee Race and Gender

```
data.support$white_female <- as.factor(data.support$treat_gender == 2 &
                                       data.support$treat_race == 1)
```

```r
data.support$black_female <-as.factor(data.support$treat_gender == 2 &
                                      data.support$treat_race == 2)
data.support$hispanic_female <- as.factor(data.support$treat_gender == 2 &
                                      data.support$treat_race == 3)
data.support$black_male <- as.factor(data.support$treat_gender == 1 &
                                      data.support$treat_race == 2)
data.support$hispanic_male <- as.factor(data.support$treat_gender == 1 &
                                      data.support$treat_race == 3)

# Overall Support
results.support.indicators <- amce(support2 ~ politicized+ treat_age + treat_lawS +
                                    treat_currentP + treat_pAbortion + white_female +
                                    black_female + hispanic_female + black_male +
                                    hispanic_male, data=data.support,cluster=TRUE,
                                    respondent.id="caseid",weights="weight")
summary(results.support.indicators)
```

```
## ------------------------------------------
## Average Marginal Component Effects (AMCE):
## ------------------------------------------
##        Attribute Level    Estimate Std. Err   z value   Pr(>|z|)
##     black_female  TRUE  0.05365689 0.023032  2.329666 1.9824e-02   *
##       black_male  TRUE  0.06598353 0.023966  2.753224 5.9011e-03  **
##  hispanic_female  TRUE -0.00202223 0.022121 -0.091415 9.2716e-01
##    hispanic_male  TRUE  0.02156142 0.023299  0.925422 3.5475e-01
##      politicized     1 -0.02112558 0.016731 -1.262650 2.0672e-01
##        treat_age     2  0.01383827 0.016585  0.834360 4.0408e-01
##        treat_age     3  0.00185229 0.016228  0.114142 9.0913e-01
##   treat_currentP     2 -0.01204190 0.020371 -0.591122 5.5444e-01
##   treat_currentP     3  0.00035078 0.021721  0.016149 9.8712e-01
##   treat_currentP     4 -0.03795573 0.020240 -1.875289 6.0753e-02
##   treat_currentP     5 -0.04798653 0.020613 -2.327984 1.9913e-02   *
##       treat_lawS     2 -0.01990847 0.018775 -1.060388 2.8897e-01
##       treat_lawS     3 -0.04184061 0.018784 -2.227440 2.5918e-02   *
##       treat_lawS     4 -0.08541911 0.018068 -4.727647 2.2714e-06 ***
##  treat_pAbortion     2 -0.09151611 0.017235 -5.309943 1.0966e-07 ***
##  treat_pAbortion     3 -0.08263842 0.018691 -4.421204 9.8152e-06 ***
##     white_female  TRUE  0.02238579 0.022583  0.991279 3.2155e-01
## ---
## Number of Obs. = 9947
## ---
## Number of Respondents = 2499
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05
##
## --------------------
## AMCE Baseline Levels:
## --------------------
##        Attribute Level
##     black_female
##       black_male
##  hispanic_female
##    hispanic_male
##      politicized
```

```
##        treat_age
##   treat_currentP
##       treat_lawS
##  treat_pAbortion
##     white_female
```

```
# Republican support
data.support.rep <- data.support[data.support$pid3a == "Republican",]
results.support.indicators.republicans <- amce(support2 ~ politicized+ treat_age +
                                                 treat_lawS + treat_currentP +
                                                 treat_pAbortion + white_female +
                                                 black_female + hispanic_female +
                                                 black_male +  hispanic_male,
                                                 data=data.support.rep,cluster=TRUE,
                                                 respondent.id="caseid",weights="weight")
summary(results.support.indicators.republicans)
```

```
## ----------------------------------------
## Average Marginal Component Effects (AMCE):
## ----------------------------------------
##        Attribute Level    Estimate Std. Err   z value   Pr(>|z|)
##     black_female  TRUE -0.01875969 0.041790 -0.448902 6.5350e-01
##       black_male  TRUE -0.04637111 0.040134 -1.155418 2.4792e-01
##  hispanic_female  TRUE -0.06407982 0.041264 -1.552929 1.2044e-01
##    hispanic_male  TRUE -0.00055369 0.038698 -0.014308 9.8858e-01
##      politicized     1  0.05722446 0.030555  1.872823 6.1093e-02
##        treat_age     2  0.01183791 0.029619  0.399675 6.8940e-01
##        treat_age     3 -0.01592992 0.032772 -0.486088 6.2690e-01
##   treat_currentP     2 -0.03412307 0.039588 -0.861955 3.8871e-01
##   treat_currentP     3  0.02328112 0.042249  0.551049 5.8160e-01
##   treat_currentP     4 -0.06960865 0.036089 -1.928788 5.3757e-02
##   treat_currentP     5 -0.02346092 0.041157 -0.570035 5.6865e-01
##       treat_lawS     2 -0.01729708 0.031907 -0.542109 5.8774e-01
##       treat_lawS     3 -0.05071579 0.035855 -1.414474 1.5722e-01
##       treat_lawS     4 -0.11015036 0.032887 -3.349334 8.1006e-04 ***
##  treat_pAbortion     2  0.02297030 0.033507  0.685534 4.9301e-01
##  treat_pAbortion     3  0.24819671 0.036260  6.844902 7.6528e-12 ***
##     white_female  TRUE -0.02244716 0.042165 -0.532363 5.9447e-01
## ---
## Number of Obs. = 2584
## ---
## Number of Respondents = 649
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05
##
## --------------------
## AMCE Baseline Levels:
## --------------------
##        Attribute Level
##     black_female
##       black_male
##  hispanic_female
##    hispanic_male
##      politicized
##        treat_age
```

```
##   treat_currentP
##       treat_lawS
##  treat_pAbortion
##     white_female
```

```
# Democrat support
data.support.dem <- data.support[data.support$pid3a == "Democrat",]
results.support.indicators.democrats <- amce(support2 ~ politicized+ treat_age +
                                               treat_lawS + treat_currentP +
                                               treat_pAbortion + white_female +
                                               black_female + hispanic_female +
                                               black_male + hispanic_male,
                                               data=data.support.dem,cluster=TRUE,
                                               respondent.id="caseid",weights="weight")
summary(results.support.indicators.democrats)
```

```
## ----------------------------------------
## Average Marginal Component Effects (AMCE):
## ----------------------------------------
##        Attribute Level   Estimate Std. Err    z value   Pr(>|z|)
##     black_female  TRUE  0.1116695 0.038650   2.889268 3.8614e-03  **
##       black_male  TRUE  0.1333526 0.045865   2.907491 3.6434e-03  **
##  hispanic_female  TRUE  0.0300189 0.037519   0.800091 4.2366e-01
##    hispanic_male  TRUE  0.0813230 0.040284   2.018756 4.3513e-02   *
##      politicized     1 -0.1221560 0.030572  -3.995700 6.4503e-05 ***
##        treat_age     2  0.0016258 0.032685   0.049741 9.6033e-01
##        treat_age     3 -0.0044093 0.027471  -0.160506 8.7248e-01
##   treat_currentP     2  0.0246288 0.031519   0.781389 4.3457e-01
##   treat_currentP     3  0.0182297 0.036023   0.506055 6.1282e-01
##   treat_currentP     4 -0.0168837 0.035103  -0.480970 6.3054e-01
##   treat_currentP     5 -0.0444733 0.034944  -1.272706 2.0312e-01
##       treat_lawS     2 -0.0318527 0.032591  -0.977342 3.2840e-01
##       treat_lawS     3 -0.0439569 0.033504  -1.311974 1.8953e-01
##       treat_lawS     4 -0.0554216 0.029130  -1.902578 5.7096e-02
##  treat_pAbortion     2 -0.2258525 0.030767  -7.340687 2.1250e-13 ***
##  treat_pAbortion     3 -0.3231943 0.031191 -10.361927 3.6944e-25 ***
##     white_female  TRUE  0.0487012 0.041626   1.169980 2.4201e-01
## ---
## Number of Obs. = 3417
## ---
## Number of Respondents = 858
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05
##
## --------------------
## AMCE Baseline Levels:
## --------------------
##        Attribute Level
##     black_female
##       black_male
##  hispanic_female
##    hispanic_male
##      politicized
##        treat_age
##   treat_currentP
```

```
##       treat_lawS
##  treat_pAbortion
##     white_female
```

```
# White Male Republican support
data.support.white.male.rep <- data.support[data.support$pid3a == "Republican" &
                                            data.support$race_respondent == "White" &
                                            data.support$gender_respondent == "Male",]
results.support.indicators.white.male.republicans <- amce(support2 ~ politicized+ treat_age +
                                                          treat_lawS + treat_currentP +
                                                          treat_pAbortion +  white_female +
                                                          black_female + hispanic_female +
                                                          black_male + hispanic_male,
                                                          data=data.support.white.male.rep,
                                                          cluster=TRUE, respondent.id="caseid",
                                                          weights="weight")
summary(results.support.indicators.white.male.republicans)
```

```
## ----------------------------------------
## Average Marginal Component Effects (AMCE):
## ----------------------------------------
##        Attribute Level   Estimate Std. Err  z value   Pr(>|z|)
##     black_female  TRUE -0.1080243 0.066462 -1.62536 1.0409e-01
##       black_male  TRUE -0.1144800 0.057328 -1.99693 4.5833e-02   *
##  hispanic_female  TRUE -0.1545313 0.065168 -2.37127 1.7727e-02   *
##    hispanic_male  TRUE -0.0702503 0.058798 -1.19478 2.3217e-01
##      politicized     1  0.1444761 0.047439  3.04554 2.3226e-03  **
##        treat_age     2  0.0099568 0.048227  0.20646 8.3643e-01
##        treat_age     3  0.0280412 0.057842  0.48479 6.2782e-01
##   treat_currentP     2 -0.0997899 0.062237 -1.60338 1.0885e-01
##   treat_currentP     3 -0.0054464 0.054208 -0.10047 9.1997e-01
##   treat_currentP     4 -0.0549426 0.056582 -0.97102 3.3154e-01
##   treat_currentP     5 -0.1007145 0.060620 -1.66139 9.6634e-02
##       treat_lawS     2  0.0390539 0.051256  0.76194 4.4610e-01
##       treat_lawS     3  0.0201417 0.055085  0.36565 7.1463e-01
##       treat_lawS     4 -0.0943310 0.049594 -1.90206 5.7164e-02
##  treat_pAbortion     2  0.0100334 0.050069  0.20039 8.4118e-01
##  treat_pAbortion     3  0.2343523 0.052689  4.44788 8.6723e-06 ***
##     white_female  TRUE -0.0220330 0.060056 -0.36687 7.1371e-01
## ---
## Number of Obs. = 1082
## ---
## Number of Respondents = 271
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05
##
## --------------------
## AMCE Baseline Levels:
## --------------------
##        Attribute Level
##     black_female
##       black_male
##  hispanic_female
##    hispanic_male
##      politicized
```

```
##        treat_age
##   treat_currentP
##       treat_lawS
##  treat_pAbortion
##     white_female
```

```
# White Male Democrat support
data.support.white.male.dem <- data.support[data.support$pid3a == "Democrat" &
                                            data.support$race_respondent == "White" &
                                            data.support$gender_respondent == "Male",]
results.support.indicators.white.male.democrats <- amce(support2 ~ politicized+ treat_age +
                                                        treat_lawS + treat_currentP +
                                                        treat_pAbortion +  white_female +
                                                        black_female +  hispanic_female +
                                                        black_male + hispanic_male,
                                                        data=data.support.white.male.dem,
                                                        cluster=TRUE, respondent.id="caseid",
                                                        weights="weight")
summary(results.support.indicators.white.male.democrats)
```

```
## ------------------------------------------
## Average Marginal Component Effects (AMCE):
## ------------------------------------------
##        Attribute Level   Estimate Std. Err   z value   Pr(>|z|)
##     black_female  TRUE  0.1143308 0.075695  1.510412 1.3094e-01
##       black_male  TRUE  0.1753180 0.093814  1.868791 6.1652e-02
##  hispanic_female  TRUE  0.0289359 0.059417  0.487002 6.2626e-01
##    hispanic_male  TRUE  0.0110065 0.059898  0.183755 8.5421e-01
##      politicized     1 -0.0194409 0.058852 -0.330338 7.4114e-01
##        treat_age     2  0.0013605 0.060299  0.022563 9.8200e-01
##        treat_age     3 -0.0434169 0.060785 -0.714268 4.7506e-01
##   treat_currentP     2  0.0413258 0.057360  0.720457 4.7124e-01
##   treat_currentP     3 -0.0174919 0.058218 -0.300455 7.6383e-01
##   treat_currentP     4 -0.1193770 0.068076 -1.753574 7.9504e-02
##   treat_currentP     5 -0.0914974 0.066918 -1.367313 1.7153e-01
##       treat_lawS     2 -0.0148368 0.059713 -0.248469 8.0377e-01
##       treat_lawS     3 -0.0676833 0.053735 -1.259586 2.0782e-01
##       treat_lawS     4 -0.0132233 0.046907 -0.281906 7.7802e-01
##  treat_pAbortion     2 -0.2266755 0.067525 -3.356903 7.8821e-04 ***
##  treat_pAbortion     3 -0.3648903 0.065711 -5.552981 2.8084e-08 ***
##     white_female  TRUE  0.0891934 0.076055  1.172754 2.4089e-01
## ---
## Number of Obs. = 909
## ---
## Number of Respondents = 228
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05
##
## --------------------
## AMCE Baseline Levels:
## --------------------
##        Attribute Level
##     black_female
##       black_male
##  hispanic_female
```

```
##    hispanic_male
##      politicized
##        treat_age
##   treat_currentP
##       treat_lawS
##  treat_pAbortion
##     white_female
```

```
# White Female Democrat support
data.support.white.female.dem <- data.support[data.support$pid3a == "Democrat" &
                                                data.support$race_respondent == "White" &
                                                data.support$gender_respondent == "Female",]
results.support.indicators.white.female.democrats <- amce(support2 ~ politicized+ treat_age +
                                                            treat_lawS +  treat_currentP +
                                                            treat_pAbortion +  white_female +
                                                            black_female + hispanic_female +
                                                            black_male +  hispanic_male,
                                                            data=data.support.white.female.dem,
                                                            cluster=TRUE,respondent.id="caseid",
                                                            weights="weight")
summary(results.support.indicators.white.female.democrats)
```

```
## ----------------------------------------
## Average Marginal Component Effects (AMCE):
## ----------------------------------------
##        Attribute Level   Estimate Std. Err   z value   Pr(>|z|)
##     black_female  TRUE  0.0220728 0.051675   0.42715 6.6927e-01
##       black_male  TRUE  0.0309415 0.062212   0.49735 6.1894e-01
##  hispanic_female  TRUE  0.0790830 0.062108   1.27331 2.0291e-01
##    hispanic_male  TRUE  0.0871769 0.052709   1.65392 9.8145e-02
##      politicized     1 -0.2193449 0.035850  -6.11843 9.4503e-10 ***
##        treat_age     2  0.0280706 0.043429   0.64635 5.1805e-01
##        treat_age     3  0.0261062 0.034746   0.75134 4.5245e-01
##   treat_currentP     2  0.0081403 0.048969   0.16623 8.6797e-01
##   treat_currentP     3 -0.0508477 0.045170  -1.12569 2.6029e-01
##   treat_currentP     4 -0.0539848 0.052497  -1.02835 3.0379e-01
##   treat_currentP     5 -0.0739161 0.048274  -1.53117 1.2573e-01
##       treat_lawS     2 -0.0549290 0.043772  -1.25489 2.0952e-01
##       treat_lawS     3 -0.0239863 0.040958  -0.58563 5.5813e-01
##       treat_lawS     4 -0.0942125 0.037839  -2.48985 1.2780e-02   *
##  treat_pAbortion     2 -0.3166364 0.043128  -7.34183 2.1069e-13 ***
##  treat_pAbortion     3 -0.4620708 0.033618 -13.74492 5.4632e-43 ***
##     white_female  TRUE  0.0482418 0.051777   0.93172 3.5148e-01
## ---
## Number of Obs. = 1371
## ---
## Number of Respondents = 344
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05
##
## --------------------
## AMCE Baseline Levels:
## --------------------
##        Attribute Level
##     black_female
```

```
##       black_male
##  hispanic_female
##    hispanic_male
##      politicized
##        treat_age
##   treat_currentP
##       treat_lawS
##  treat_pAbortion
##     white_female
```

```r
# White Female Republican support
data.support.white.female.rep <- data.support[data.support$pid3a == "Republican" &
                                                data.support$race_respondent == "White" &
                                                data.support$gender_respondent == "Female",]
results.support.indicators.white.female.republicans <- amce(support2 ~ politicized+ treat_age +
                                                               treat_lawS + treat_currentP +
                                                               treat_pAbortion +  white_female +
                                                               black_female + hispanic_female +
                                                               black_male + hispanic_male,
                                                               data=data.support.white.female.rep,
                                                               cluster=TRUE, respondent.id="caseid",
                                                               weights="weight")
summary(results.support.indicators.white.female.republicans)
```

```
## ------------------------------------------
## Average Marginal Component Effects (AMCE):
## ------------------------------------------
##        Attribute Level   Estimate Std. Err   z value   Pr(>|z|)
##     black_female  TRUE -0.0282203 0.047567 -0.593274 5.5300e-01
##       black_male  TRUE -0.0798188 0.056303 -1.417662 1.5629e-01
##  hispanic_female  TRUE -0.0267608 0.059405 -0.450485 6.5236e-01
##    hispanic_male  TRUE  0.0020432 0.054946  0.037186 9.7034e-01
##      politicized     1  0.0195622 0.042798  0.457076 6.4762e-01
##        treat_age     2 -0.0106512 0.039869 -0.267157 7.8935e-01
##        treat_age     3 -0.0334418 0.043100 -0.775918 4.3780e-01
##   treat_currentP     2  0.0591859 0.052206  1.133690 2.5692e-01
##   treat_currentP     3  0.0863394 0.062857  1.373583 1.6957e-01
##   treat_currentP     4 -0.0764455 0.048270 -1.583708 1.1326e-01
##   treat_currentP     5  0.0215346 0.057791  0.372632 7.0942e-01
##       treat_lawS     2 -0.0579692 0.042881 -1.351851 1.7642e-01
##       treat_lawS     3 -0.1080273 0.047724 -2.263603 2.3599e-02   *
##       treat_lawS     4 -0.0969370 0.043642 -2.221193 2.6338e-02   *
##  treat_pAbortion     2  0.0051913 0.045677  0.113653 9.0951e-01
##  treat_pAbortion     3  0.2735454 0.053418  5.120825 3.0420e-07 ***
##     white_female  TRUE -0.0480758 0.062048 -0.774815 4.3845e-01
## ---
## Number of Obs. = 1226
## ---
## Number of Respondents = 308
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05
##
## --------------------
## AMCE Baseline Levels:
## --------------------
```

```
##        Attribute Level
##     black_female
##       black_male
##  hispanic_female
##    hispanic_male
##      politicized
##        treat_age
##   treat_currentP
##       treat_lawS
##  treat_pAbortion
##     white_female
```

Table B.1: Descriptive Statistics

```
round(prop.table(table(data$gender_respondent)), 3)
```

```
##
##         Male       Female   [8]skipped [9]not asked
##        0.465        0.535        0.000        0.000
```

```
round(prop.table(table(data$race_respondent)), 3)
```

```
##
##           White           Black        Hispanic           Asian
##           0.773           0.092           0.078           0.020
## Native American           Mixed           Other  Middle Eastern
##           0.006           0.018           0.008           0.004
##     [98]skipped   [99]not asked
##           0.000           0.000
```

```
round(prop.table(table(data$educ)), 3)
```

```
##
##                No HS High school graduate         Some college
##                0.042                0.361               0.228
##               2-year               4-year            Post-grad
##                0.095                0.180               0.094
##           [8]skipped         [9]not asked
##                0.000                0.000
```

```
round(prop.table(table(data$faminc2)), 3)
```

```
##
## $100,000 to $149,999     $150,000 or more   $20,000 to $39,999
##                0.075                0.036               0.226
##   $40,000 to $59,999   $60,000 to $79,999   $80,000 to $99,999
##                0.164                0.118               0.071
##    Prefer not to say         Under 20,000
##                0.129                0.180
```

```
round(prop.table(table(data$pid3)), 3)
```

```
##
##     Democrat   Republican  Independent        Other     Not sure
##        0.344        0.260        0.299        0.045        0.053
```

```
##   [8]skipped [9]not asked
##        0.000        0.000
```

```
round(prop.table(table(data$ideo5)), 3)
```

```
##
##      Very liberal           Liberal          Moderate      Conservative
##             0.096             0.186             0.328             0.240
## Very conservative         Not sure       [8]skipped      [9]not asked
##             0.072             0.076             0.000             0.000
```

Table B.3: Effect of Descriptive Representation on Attitudes toward Judicial Nominees: Full Results

```
# Estimate the AMCEs for support (binary), cluster SEs by respondent
results.support.overall.binary <- amce(support2 ~ politicized + treat_age + treat_gender + treat_race +
                                  same.race, data=data.support,cluster=TRUE, respondent.id="caseid", we
summary(results.support.overall.binary)
```

```
## ------------------------------------------
## Average Marginal Component Effects (AMCE):
## ------------------------------------------
##        Attribute Level    Estimate Std. Err   z value   Pr(>|z|)
##      politicized     1 -0.02152808 0.016788 -1.282321 1.9973e-01
##      same.gender     1  0.01267180 0.012395  1.022313 3.0663e-01
##        same.race     1  0.06145415 0.017882  3.436691 5.8887e-04 ***
##        treat_age     2  0.01262425 0.016684  0.756663 4.4925e-01
##        treat_age     3  0.00218115 0.016258  0.134157 8.9328e-01
##   treat_currentP     2 -0.01297930 0.020336 -0.638240 5.2332e-01
##   treat_currentP     3  0.00022289 0.021704  0.010269 9.9181e-01
##   treat_currentP     4 -0.03876031 0.020154 -1.923231 5.4451e-02
##   treat_currentP     5 -0.04926600 0.020614 -2.389920 1.6852e-02   *
##     treat_gender     2 -0.00518206 0.012479 -0.415263 6.7795e-01
##       treat_lawS     2 -0.01972457 0.018764 -1.051194 2.9317e-01
##       treat_lawS     3 -0.04386852 0.018859 -2.326152 2.0010e-02   *
##       treat_lawS     4 -0.08701392 0.018071 -4.815162 1.4708e-06 ***
##  treat_pAbortion     2 -0.09254834 0.017174 -5.388778 7.0938e-08 ***
##  treat_pAbortion     3 -0.08346417 0.018725 -4.457367 8.2973e-06 ***
##       treat_race     2  0.08244811 0.018797  4.386291 1.1530e-05 ***
##       treat_race     3  0.03215455 0.018638  1.725198 8.4492e-02
## ---
## Number of Obs. = 9947
## ---
## Number of Respondents = 2499
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05
##
## --------------------
## AMCE Baseline Levels:
## --------------------
##        Attribute Level
##      politicized
```

```
##      same.gender
##        same.race
##        treat_age
##   treat_currentP
##     treat_gender
##       treat_lawS
##  treat_pAbortion
##       treat_race
```

```
# Estimate the AMCEs trust (binary), cluster SEs by respondent
results.trust.overall.binary <- amce(trust2 ~ politicized + treat_age + treat_gender + treat_race +  tr
                                same.race, data=data.trust,cluster=TRUE, respondent.id="caseid", weights
summary(results.trust.overall.binary)
```

```
## ----------------------------------------
## Average Marginal Component Effects (AMCE):
## ----------------------------------------
##        Attribute Level   Estimate Std. Err  z value   Pr(>|z|)
##      politicized     1  0.0097007 0.017415  0.55703 5.7751e-01
##      same.gender     1  0.0249145 0.012216  2.03957 4.1393e-02   *
##        same.race     1  0.0622936 0.020010  3.11314 1.8511e-03  **
##        treat_age     2 -0.0049870 0.015419 -0.32343 7.4637e-01
##        treat_age     3 -0.0082465 0.015757 -0.52335 6.0073e-01
##   treat_currentP     2 -0.0568071 0.021775 -2.60877 9.0868e-03  **
##   treat_currentP     3 -0.0205363 0.020944 -0.98054 3.2682e-01
##   treat_currentP     4 -0.0662148 0.020251 -3.26978 1.0763e-03  **
##   treat_currentP     5 -0.0794167 0.020276 -3.91671 8.9766e-05 ***
##     treat_gender     2 -0.0098238 0.012247 -0.80212 4.2248e-01
##       treat_lawS     2 -0.0102141 0.017067 -0.59848 5.4952e-01
##       treat_lawS     3 -0.0270695 0.017549 -1.54249 1.2295e-01
##       treat_lawS     4 -0.0440743 0.017696 -2.49066 1.2751e-02   *
##  treat_pAbortion     2 -0.0498379 0.017477 -2.85158 4.3502e-03  **
##  treat_pAbortion     3 -0.0662886 0.017824 -3.71910 1.9994e-04 ***
##       treat_race     2  0.0695487 0.019811  3.51057 4.4714e-04 ***
##       treat_race     3  0.0501062 0.020020  2.50275 1.2323e-02   *
## ---
## Number of Obs. = 9970
## ---
## Number of Respondents = 2500
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05
##
## --------------------
## AMCE Baseline Levels:
## --------------------
##        Attribute Level
##      politicized
##      same.gender
##        same.race
##        treat_age
##   treat_currentP
##     treat_gender
##       treat_lawS
##  treat_pAbortion
##       treat_race
```

```r
# Estimate the AMCEs qualified (binary), cluster SEs by respondent
results.qualified.overall.binary <- amce(qualified2 ~ politicized + treat_age + treat_gender + treat_ra
                                         same.race, data=data.qualified,cluster=TRUE, respondent.id="caseid"
summary(results.qualified.overall.binary)
```

```
## ----------------------------------------
## Average Marginal Component Effects (AMCE):
## ----------------------------------------
##        Attribute Level   Estimate Std. Err  z value   Pr(>|z|)
##      politicized     1 -0.0414005 0.019271 -2.14836 3.1686e-02   *
##      same.gender     1  0.0032480 0.013705  0.23700 8.1266e-01
##        same.race     1  0.0452344 0.019436  2.32730 1.9949e-02   *
##        treat_age     2  0.0025042 0.016949  0.14775 8.8254e-01
##        treat_age     3 -0.0209272 0.016512 -1.26738 2.0502e-01
##   treat_currentP     2 -0.1069151 0.022754 -4.69866 2.6188e-06 ***
##   treat_currentP     3 -0.0444020 0.024187 -1.83577 6.6392e-02
##   treat_currentP     4 -0.1356271 0.023242 -5.83535 5.3678e-09 ***
##   treat_currentP     5 -0.1508390 0.023122 -6.52356 6.8659e-11 ***
##     treat_gender     2  0.0050808 0.013729  0.37007 7.1133e-01
##       treat_lawS     2 -0.0397186 0.020082 -1.97781 4.7950e-02   *
##       treat_lawS     3 -0.0670809 0.019506 -3.43906 5.8375e-04 ***
##       treat_lawS     4 -0.1557614 0.018804 -8.28347 1.1964e-16 ***
##  treat_pAbortion     2 -0.0509654 0.018363 -2.77538 5.5137e-03  **
##  treat_pAbortion     3 -0.0458375 0.018416 -2.48899 1.2811e-02   *
##       treat_race     2  0.0642285 0.019598  3.27734 1.0479e-03  **
##       treat_race     3  0.0301445 0.020199  1.49235 1.3561e-01
## ---
## Number of Obs. = 9955
## ---
## Number of Respondents = 2499
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05
##
## --------------------
## AMCE Baseline Levels:
## --------------------
##        Attribute Level
##      politicized
##      same.gender
##        same.race
##        treat_age
##   treat_currentP
##     treat_gender
##       treat_lawS
##  treat_pAbortion
##       treat_race
```

Figure C.1: Effect of Shared Descriptive Characteristics on Nominee Evaluations (Bootstrapped Standard Errors)

```
# Set the seed to run the bootstrap and create empty lists to store results
set.seed(02138)
nboot <- 1000

supportresults  <- list()
qualifiedresults  <- list()
trustresults  <- list()

# Unique respondent IDs
caseids.support <- unique(data.support$caseid)
caseids.qualified <- unique(data.qualified$caseid)
caseids.trust <- unique(data.trust$caseid)

for (k in 1:nboot) {

  # Sample respondents to be used in this bootstrap iteration
  respondent_sample_support   <- sample(caseids.support,  length(caseids.support),replace=TRUE)
  respondent_sample_qualified <- sample(caseids.qualified,length(caseids.qualified),replace=TRUE)
  respondent_sample_trust     <- sample(caseids.trust,    length(caseids.trust),replace=TRUE)

  # Sample all of the profiles that respondent received
  support_dataset <-   inner_join(tibble(caseid = respondent_sample_support),
                                  data.support, by = "caseid")
  qualified_dataset <- inner_join(tibble(caseid = respondent_sample_qualified),
                                  data.qualified, by = "caseid")
  trust_dataset <-     inner_join(tibble(caseid = respondent_sample_trust),
                                  data.trust, by = "caseid")

  # Then, run regressions
  support_bootdat_nocov <- amce(support2 ~ politicized + treat_age + treat_gender +
                                  treat_race +  treat_lawS + treat_currentP +
                                  treat_pAbortion + same.gender + same.race,
                                  data=support_dataset,cluster=TRUE,
                                  respondent.id="caseid",weights="weight")

  qualified_bootdat_nocov <- amce(qualified2 ~ politicized + treat_age + treat_gender +
                                    treat_race +  treat_lawS + treat_currentP +
                                    treat_pAbortion + same.gender + same.race,
                                    data=qualified_dataset,cluster=TRUE,
                                    respondent.id="caseid",weights="weight")

  trust_bootdat_nocov <- amce(trust2 ~ politicized + treat_age + treat_gender +
                                treat_race +  treat_lawS + treat_currentP +
                                treat_pAbortion + same.gender + same.race,
                                data=trust_dataset,cluster=TRUE,
                                respondent.id="caseid",weights="weight")



  supportresults[[k]]  <-lapply(support_bootdat_nocov$estimates,
```

```
                              function(i) i[1])[c("samerace", "samegender")]
  qualifiedresults[[k]]  <-lapply(qualified_bootdat_nocov$estimates,
                              function(i) i[1])[c("samerace", "samegender")]
  trustresults[[k]]  <-lapply(trust_bootdat_nocov$estimates,
                              function(i) i[1])[c("samerace", "samegender")]

  if (k %% 250 == 0) cat(paste(k, "out of", nboot, "samples stored..\n"))

}
```

```
## 250 out of 1000 samples stored..
## 500 out of 1000 samples stored..
## 750 out of 1000 samples stored..
## 1000 out of 1000 samples stored..
```

```
# Turn the bootstrap results into our CI estimates


# SEs
support_vec_se    <- apply(do.call(rbind.data.frame,supportresults), 2, sd)
qualified_vec_se  <- apply(do.call(rbind.data.frame,qualifiedresults), 2, sd)
trust_vec_se      <- apply(do.call(rbind.data.frame,trustresults), 2, sd)

# point estimates
support_vec_est    <- apply(do.call(rbind.data.frame,supportresults), 2, mean)
qualified_vec_est  <- apply(do.call(rbind.data.frame,qualifiedresults), 2, mean)
trust_vec_est     <- apply(do.call(rbind.data.frame,trustresults), 2, mean)



# Data frames for plotting
gender_support <- data.frame(est = support_vec_est["samegender"],
                             lb =  support_vec_est["samegender"] -
                             qnorm(.975)*support_vec_se["samegender"],
                             ub =  support_vec_est["samegender"] +
                             qnorm(.975)*support_vec_se["samegender"],
                             model = "Support")


race_support <-  data.frame(est = support_vec_est["samerace"],
                            lb =  support_vec_est["samerace"] -
                            qnorm(.975)*support_vec_se["samerace"],
                            ub =  support_vec_est["samerace"] +
                            qnorm(.975)*support_vec_se["samerace"],
                            model = "Support")

gender_qualified <- data.frame(est = qualified_vec_est["samegender"],
                               lb  = qualified_vec_est["samegender"] -
                               qnorm(.975)*qualified_vec_se["samegender"],
                               ub  = qualified_vec_est["samegender"] +
                               qnorm(.975)*qualified_vec_se["samegender"],
                               model = "Qualified")
```

```r
race_qualified <-  data.frame(est = qualified_vec_est["samerace"],
                              lb  = qualified_vec_est["samerace"] -
                              qnorm(.975)*qualified_vec_se["samerace"],
                              ub  = qualified_vec_est["samerace"] +
                              qnorm(.975)*qualified_vec_se["samerace"],
                              model = "Qualified")


gender_trust <- data.frame(est = trust_vec_est["samegender"],
                           lb  = trust_vec_est["samegender"] -
                           qnorm(.975)*trust_vec_se["samegender"],
                           ub  = trust_vec_est["samegender"] +
                           qnorm(.975)*trust_vec_se["samegender"],
                           model = "Trust")


race_trust <-  data.frame(est = trust_vec_est["samerace"],
                          lb  = trust_vec_est["samerace"] -
                          qnorm(.975)*trust_vec_se["samerace"],
                          ub  = trust_vec_est["samerace"] +
                          qnorm(.975)*trust_vec_se["samerace"],
                          model = "Trust")

dat <- as.data.frame(c("Qualified","Trust","Support"))
colnames(dat) <- c("order")

# Gender
p1 <-ggplot(gender_trust, aes(x = model, y = est)) +
  geom_point() +
  geom_errorbar(aes(ymin = lb, ymax = ub), width = 0.2) +
  geom_hline(yintercept = 0, lty = 2) +
  xlab("") +
  ylab("Coefficient Estimate") +
  geom_point(data= gender_qualified, aes(x = model, y = est)) +
  geom_errorbar(data=gender_qualified, aes(ymin = lb, ymax = ub), width = 0.2) +
  geom_point(data= gender_support, aes(x = model, y = est)) +
  geom_errorbar(data=gender_support, aes(ymin = lb, ymax = ub), width = 0.2) +
  theme_bw() + coord_flip() + labs(title="Gender")  + ylim(-.1,0.2) +
  scale_x_discrete(limits=dat$order) +
  theme(plot.title = element_text(hjust = 0.5))


# Race
p2 <- ggplot(race_trust, aes(x = model, y = est)) +
  geom_point() +
  geom_errorbar(aes(ymin = lb, ymax = ub), width = 0.2) +
  geom_hline(yintercept = 0, lty = 2) +
  xlab("") +
  ylab("Coefficient Estimate") +
  geom_point(data= race_qualified, aes(x = model, y = est)) +
  geom_errorbar(data=race_qualified, aes(ymin = lb, ymax = ub), width = 0.2) +
  geom_point(data= race_support, aes(x = model, y = est)) +
  geom_errorbar(data=race_support, aes(ymin = lb, ymax = ub), width = 0.2) +
```

```
  theme_bw() + coord_flip() + labs(title="Race")  + ylim(-.1,0.2) +
  scale_x_discrete(limits=dat$order) +
  theme(plot.title = element_text(hjust = 0.5))

grid.arrange(p2, p1, ncol=2, nrow=1)
```

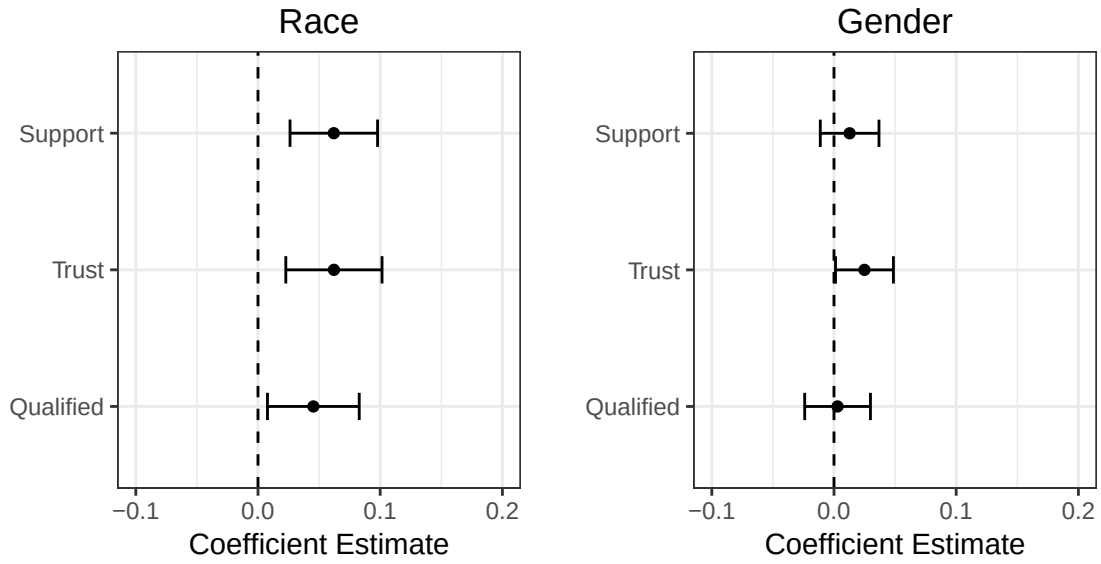


Figure C.2: Partisanship, Race, and Nominee Support (Bootstrapped Standard Errors)

```
# Create empty lists to store bootstrap results
supportresults.whitedems  <-list()
supportresults.whitereps  <-list()
supportresults.blackdems  <-list()
supportresults.blackreps  <-list()
supportresults.hispanicdems  <-list()
supportresults.hispanicreps <- list()
supportresults.womendems <- list()
supportresults.womenreps <- list()
supportresults.mendems <- list()
supportresults.menreps <- list()
ran.black.rep.regression <- rep(NA, nboot)

for(k in 1:nboot){

  # Sample respondents to be used in this bootstrap iteration

  respondent_sample_support   <- sample(caseids.support,  length(caseids.support),replace=TRUE)
  support_dataset <-   inner_join(tibble(caseid = respondent_sample_support),
                                  data.support, by = "caseid")

  # Then, run regressions #

  # Re-level the data to make sure that we are using respondent's race as the baseline
```

```r
support_dataset <- within(support_dataset, race_nominee <- relevel(race_nominee,
                                                                   ref = "White"))
support_dataset <- within(support_dataset, gender_nominee <- relevel(gender_nominee,
                                                                   ref = "Male"))


# Support, white respondents
support_dataset.white <- support_dataset[which(support_dataset$race_respondent ==
                                               "White"),]

# Democrats only
support_dataset.white.dems <- support_dataset.white[which(support_dataset.white$pid3a ==
                                                          "Democrat"),]
results.support.white.dems <- amce(support2 ~ gender_nominee + race_nominee +
                                     politicized + treat_age + treat_lawS +
                                     treat_currentP + treat_pAbortion + same.gender,
                                     data=support_dataset.white.dems,cluster=TRUE,
                                     respondent.id="caseid",weights="weight")

# Republicans only
support_dataset.white.reps <- support_dataset.white[which(support_dataset.white$pid3a ==
                                                          "Republican"),]
results.support.white.reps <- amce(support2 ~ gender_nominee + race_nominee + politicized +
                                   treat_age +  treat_lawS + treat_currentP +
                                   treat_pAbortion + same.gender,
                                   data=support_dataset.white.reps,cluster=TRUE,
                                   respondent.id="caseid",weights="weight")

# Support, Black respondents

# Re-level the data to make sure that we are using respondent's race as the baseline
support_dataset <- within(support_dataset, race_nominee <- relevel(race_nominee, ref =
                                                                     "Black"))
support_dataset.black <- support_dataset[which(support_dataset$race_respondent ==
                                               "Black"),]


# Democrats only
support_dataset.black.dems <- support_dataset.black[which(support_dataset.black$pid3a ==
                                                          "Democrat"),]
results.support.black.dems <- amce(support2 ~ gender_nominee + race_nominee +
                                     politicized + treat_age + treat_lawS +
                                     treat_currentP + treat_pAbortion + same.gender,
                                     data=support_dataset.black.dems,cluster=TRUE,
                                     respondent.id="caseid",weights="weight")

# Republicans only
support_dataset.black.reps <- support_dataset.black[which(support_dataset.black$pid3a ==
                                                          "Republican"),]

# Ensuring there is enough variation in the data to run the regression. This may occur
# due to low number of black Reps in sample and respondnets receiving the same value of
# politicized across all four profiles
```

```r
if (length(unique(support_dataset.black.reps$politicized)) > 1){
  results.support.black.reps <- amce(support2 ~ gender_nominee + race_nominee +
                                       politicized + treat_age + treat_lawS +
                                       treat_currentP + treat_pAbortion + same.gender,
                                       data=support_dataset.black.reps,cluster=TRUE,
                                       respondent.id="caseid",weights="weight")
  ran.black.rep.regression[k] <- 1
}

# Support, Hispanic respondents

# Re-level the data to make sure that we are using respondent's race as the baseline
support_dataset <- within(support_dataset, race_nominee <- relevel(race_nominee,
                                                                   ref = "Hispanic"))

support_dataset.hispanic <- support_dataset[which(support_dataset$race_respondent ==
                                                    "Hispanic"),]

# Democrats only
support_dataset.hispanic.dems <- support_dataset.hispanic[which(support_dataset.hispanic$pid3a ==
                                                                  "Democrat"),]
results.support.hispanic.dems <- amce(support2 ~ gender_nominee + race_nominee +
                                       politicized + treat_age + treat_lawS +
                                       treat_currentP + treat_pAbortion + same.gender,
                                       data=support_dataset.hispanic.dems,cluster=TRUE,
                                       respondent.id="caseid",weights="weight")

# Republicans only
support_dataset.hispanic.reps <- support_dataset.hispanic[which(support_dataset.hispanic$pid3a ==
                                                                  "Republican"),]
results.support.hispanic.reps <- amce(support2 ~ gender_nominee + race_nominee +
                                       politicized + treat_age +  treat_lawS +
                                       treat_currentP + treat_pAbortion + same.gender,
                                       data=support_dataset.hispanic.reps,cluster=TRUE,
                                       respondent.id="caseid",weights="weight")


# Now looking at support by gender match

support_dataset.male <- support_dataset[which(support_dataset$gender_respondent ==
                                                "Male"),]

# Democrats only
support_dataset.male.dems <- support_dataset.male[which(support_dataset.male$pid3a ==
                                                          "Democrat"),]
results.support.male.dems <- amce(support2 ~ gender_nominee + race_nominee +
                                    politicized + treat_age + treat_lawS +
                                    treat_currentP + treat_pAbortion + same.race,
                                    data=support_dataset.male.dems,cluster=TRUE,
                                    respondent.id="caseid",weights="weight")

# Republicans only
support_dataset.male.reps <- support_dataset.male[which(support_dataset.male$pid3a ==
```

```
                                                        "Republican"),]
results.support.male.reps <- amce(support2 ~ gender_nominee + race_nominee +
                                  politicized + treat_age +  treat_lawS +
                                  treat_currentP + treat_pAbortion + same.race,
                                  data=support_dataset.male.reps,cluster=TRUE,
                                  respondent.id="caseid",weights="weight")


# Re-level the data to make sure that we are using respondent's gender as the baseline
support_dataset <- within(support_dataset, gender_nominee <- relevel(gender_nominee,
                                                                     ref = "Female"))
support_dataset.female <- support_dataset[which(support_dataset$gender_respondent ==
                                                 "Female"),]

# Democrats only
support_dataset.female.dems <- support_dataset.female[which(support_dataset.female$pid3a ==
                                                             "Democrat"),]
results.support.female.dems <- amce(support2 ~ gender_nominee + race_nominee +
                                    politicized + treat_age + treat_lawS +
                                    treat_currentP + treat_pAbortion + same.race,
                                    data=support_dataset.female.dems,cluster=TRUE,
                                    respondent.id="caseid",weights="weight")

# Republicans only
support_dataset.female.reps <- support_dataset.female[which(support_dataset.female$pid3a ==
                                                             "Republican"),]
results.support.female.reps <- amce(support2 ~ gender_nominee + race_nominee +
                                    politicized + treat_age + treat_lawS +
                                    treat_currentP + treat_pAbortion + same.race,
                                    data=support_dataset.female.reps,cluster=TRUE,
                                    respondent.id="caseid",weights="weight")




supportresults.whitedems[[k]]  <-lapply(results.support.white.dems$estimates,
                                    function(i) i[1,])[c("racenominee")]
supportresults.whitereps[[k]]  <-lapply(results.support.white.reps$estimates,
                                    function(i) i[1,])[c("racenominee")]
supportresults.blackdems[[k]]  <-lapply(results.support.black.dems$estimates,
                                    function(i) i[1,])[c("racenominee")]
supportresults.blackreps[[k]]  <-lapply(results.support.black.reps$estimates,
                                    function(i) i[1,])[c("racenominee")]
supportresults.hispanicdems[[k]]  <-lapply(results.support.hispanic.dems$estimates,
                                    function(i) i[1,])[c("racenominee")]
supportresults.hispanicreps[[k]]  <-lapply(results.support.hispanic.reps$estimates,
                                    function(i) i[1,])[c("racenominee")]
supportresults.womendems[[k]]  <-lapply(results.support.female.dems$estimates,
                                    function(i) i[1])[c("gendernominee")]
supportresults.womenreps[[k]]  <-lapply(results.support.female.reps$estimates,
                                    function(i) i[1])[c("gendernominee")]
supportresults.mendems[[k]]  <-lapply(results.support.male.dems$estimates,
                                    function(i) i[1])[c("gendernominee")]
```

```
  supportresults.menreps[[k]]  <-lapply(results.support.male.reps$estimates,
                                      function(i) i[1])[c("gendernominee")]

  if (k %% 250 == 0) cat(paste(k, "out of", nboot, "samples stored..\n"))

}
```

```
## 250 out of 1000 samples stored..
## 500 out of 1000 samples stored..
## 750 out of 1000 samples stored..
## 1000 out of 1000 samples stored..
```

```
# Removing iterations for which no black Republican regression could be run
supportresults.blackreps2 <- supportresults.blackreps[!is.na(ran.black.rep.regression)]

# Turn the bootstrap results into our CI estimates
# SEs
supportresults.whitedems_vec_se  <- apply(t(rbind.data.frame(supportresults.whitedems)),
                                          2, sd, na.rm=TRUE)
supportresults.whitereps_vec_se  <- apply(t(rbind.data.frame(supportresults.whitereps)),
                                          2, sd, na.rm=TRUE)
supportresults.blackdems_vec_se  <- apply(t(rbind.data.frame(supportresults.blackdems)),
                                          2, sd, na.rm=TRUE)
supportresults.blackreps_vec_se  <- apply(t(rbind.data.frame(supportresults.blackreps2)),
                                          2, sd, na.rm=TRUE)
supportresults.hispanicdems_vec_se<- apply(t(rbind.data.frame(supportresults.hispanicdems)),
                                          2, sd, na.rm=TRUE)
supportresults.hispanicreps_vec_se<- apply(t(rbind.data.frame(supportresults.hispanicreps)),
                                          2, sd, na.rm=TRUE)
supportresults.womendems_vec_se <- apply(do.call(rbind.data.frame,supportresults.womendems),
                                          2, sd, na.rm=TRUE)
supportresults.womenreps_vec_se <-  apply(do.call(rbind.data.frame,supportresults.womenreps),
                                          2, sd, na.rm=TRUE)
supportresults.mendems_vec_se   <-  apply(do.call(rbind.data.frame,supportresults.mendems),
                                          2, sd, na.rm=TRUE)
supportresults.menreps_vec_se   <-  apply(do.call(rbind.data.frame,supportresults.menreps),
                                          2, sd, na.rm=TRUE)

# Point estimates
supportresults.whitedems_vec_est  <- apply(t(rbind.data.frame(supportresults.whitedems)),
                                           2, mean, na.rm=TRUE)
supportresults.whitereps_vec_est  <- apply(t(rbind.data.frame(supportresults.whitereps)),
                                           2, mean, na.rm=TRUE)
supportresults.blackdems_vec_est  <- apply(t(rbind.data.frame(supportresults.blackdems)),
                                           2, mean, na.rm=TRUE)
supportresults.blackreps_vec_est  <- apply(t(rbind.data.frame(supportresults.blackreps2)),
                                           2, mean, na.rm=TRUE)
supportresults.hispanicdems_vec_est <- apply(t(rbind.data.frame(supportresults.hispanicdems)),
                                           2, mean, na.rm=TRUE)
supportresults.hispanicreps_vec_est <- apply(t(rbind.data.frame(supportresults.hispanicreps)),
                                           2, mean, na.rm=TRUE)
supportresults.womendems_vec_est<- apply(do.call(rbind.data.frame,supportresults.womendems),
                                           2, mean, na.rm=TRUE)
supportresults.womenreps_vec_est<-  apply(do.call(rbind.data.frame,supportresults.womenreps),
```

```
                                        2, mean, na.rm=TRUE)
supportresults.mendems_vec_est <-  apply(do.call(rbind.data.frame,supportresults.mendems),
                                        2, mean, na.rm=TRUE)
supportresults.menreps_vec_est  <-  apply(do.call(rbind.data.frame,supportresults.menreps),
                                        2, mean, na.rm=TRUE)

# Data frames for plotting
whiteresp.dem.hispanic <-  data.frame(est = supportresults.whitedems_vec_est["racenomineeHispanic"],
                                      lb = supportresults.whitedems_vec_est["racenomineeHispanic"] -  qnorm
                                      ub = supportresults.whitedems_vec_est["racenomineeHispanic"] +  qnorm
                                      model = "White-Hispanic")

whiteresp.dem.black <-  data.frame(est = supportresults.whitedems_vec_est["racenomineeBlack"],
                                       lb = supportresults.whitedems_vec_est["racenomineeBlack"] -  qnorm
                                       ub = supportresults.whitedems_vec_est["racenomineeBlack"] +  qnorm
                                       model = "White-Black")


blackresp.dem.hispanic <- data.frame(est = supportresults.blackdems_vec_est["racenomineeHispanic"],
                                     lb = supportresults.blackdems_vec_est["racenomineeHispanic"] -  qn
                                     ub = supportresults.blackdems_vec_est["racenomineeHispanic"] +  qn
                                     model = "Black-Hispanic")


blackresp.dem.white <- data.frame(est = supportresults.blackdems_vec_est["racenomineeWhite"],
                                  lb = supportresults.blackdems_vec_est["racenomineeWhite"] -  qnorm(.9
                                  ub = supportresults.blackdems_vec_est["racenomineeWhite"] +  qnorm(.9
                                  model = "Black-White")


hispanicresp.dem.white <- data.frame(est = supportresults.hispanicdems_vec_est["racenomineeWhite"],
                                     lb =  supportresults.hispanicdems_vec_est["racenomineeWhite"] -  q
                                     ub =  supportresults.hispanicdems_vec_est["racenomineeWhite"] +  q
                                     model = "Hispanic-White")


hispanicresp.dem.black <- data.frame(est = supportresults.hispanicdems_vec_est["racenomineeBlack"],
                                     lb =  supportresults.hispanicdems_vec_est["racenomineeBlack"] - qn
                                     ub = supportresults.hispanicdems_vec_est["racenomineeBlack"] +  qn
                                     model = "Hispanic-Black")

whiteresp.rep.hispanic <-  data.frame(est = supportresults.whitereps_vec_est["racenomineeHispanic"],
                                      lb = supportresults.whitereps_vec_est["racenomineeHispanic"] -  q
                                      ub = supportresults.whitereps_vec_est["racenomineeHispanic"] +  q
                                      model = "White-Hispanic")

whiteresp.rep.black <-  data.frame(est = supportresults.whitereps_vec_est["racenomineeBlack"],
                                   lb = supportresults.whitereps_vec_est["racenomineeBlack"] -  qnorm(.
                                   ub = supportresults.whitereps_vec_est["racenomineeBlack"] +  qnorm(.
                                   model = "White-Black")


blackresp.rep.hispanic <- data.frame(est = supportresults.blackreps_vec_est["racenomineeHispanic"],
```

```
                                    lb = supportresults.blackreps_vec_est["racenomineeHispanic"] -  qn
                                    ub = supportresults.blackreps_vec_est["racenomineeHispanic"] +  qn
                                    model = "Black-Hispanic")


blackresp.rep.white <- data.frame(est = supportresults.blackreps_vec_est["racenomineeWhite"],
                                  lb = supportresults.blackreps_vec_est["racenomineeWhite"] -  qnorm(.9
                                  ub = supportresults.blackreps_vec_est["racenomineeWhite"] +  qnorm(.9
                                  model = "Black-White")


hispanicresp.rep.white <- data.frame(est = supportresults.hispanicreps_vec_est["racenomineeWhite"],
                                    lb =  supportresults.hispanicreps_vec_est["racenomineeWhite"] -  q
                                    ub =  supportresults.hispanicreps_vec_est["racenomineeWhite"] +  q
                                    model = "Hispanic-White")


hispanicresp.rep.black <- data.frame(est = supportresults.hispanicreps_vec_est["racenomineeBlack"],
                                    lb =  supportresults.hispanicreps_vec_est["racenomineeBlack"] - qn
                                    ub = supportresults.hispanicreps_vec_est["racenomineeBlack"] +  qn
                                    model = "Hispanic-Black")


g3 <- ggplot(hispanicresp.dem.white, aes(x = model, y = est)) +
  geom_point(colour="grey45") +
  geom_errorbar(aes(ymin = lb, ymax = ub), width = 0.2, colour="grey45") +
  geom_hline(yintercept = 0, lty = 2) +
  xlab("") +
  ylab("") +
  geom_point(data= hispanicresp.dem.black, aes(x = model, y = est),colour="grey45") +
  geom_errorbar(data=hispanicresp.dem.black, aes(ymin = lb, ymax = ub), width = 0.2,
  colour="grey45") +
  geom_point(data= blackresp.dem.hispanic, aes(x = model, y = est),colour="grey70") +
  geom_errorbar(data=blackresp.dem.hispanic, aes(ymin = lb, ymax = ub), width = 0.2,
  colour="grey70") +
  geom_point(data= blackresp.dem.white, aes(x = model, y = est),colour="grey70") +
  geom_errorbar(data=blackresp.dem.white, aes(ymin = lb, ymax = ub), width = 0.2,
  colour="grey70")  +
  geom_point(data= whiteresp.dem.hispanic, aes(x = model, y = est),colour="black") +
  geom_errorbar(data=whiteresp.dem.hispanic, aes(ymin = lb, ymax = ub), width = 0.2,
  colour="black") +
  geom_point(data= whiteresp.dem.black, aes(x = model, y = est),colour="black") +
  geom_errorbar(data=whiteresp.dem.black, aes(ymin = lb, ymax = ub), width = 0.2,
  colour="black")  +
  theme_bw() + coord_flip() + ylim(-1.75,1.5) +
  labs(title="Democrats: Support for Nominees by Race") +
  theme(plot.title = element_text(hjust = 0.5))


g4 <- ggplot(hispanicresp.rep.white, aes(x = model, y = est)) +
  geom_point(colour="grey45") +
  geom_errorbar(aes(ymin = lb, ymax = ub), width = 0.2, colour="grey45") +
  geom_hline(yintercept = 0, lty = 2) +
```

```
  xlab("") +
  ylab("") +
  geom_point(data= hispanicresp.rep.black, aes(x = model, y = est),colour="grey45") +
  geom_errorbar(data=hispanicresp.rep.black, aes(ymin = lb, ymax = ub), width = 0.2,
  colour="grey45") +
  geom_point(data= blackresp.rep.hispanic, aes(x = model, y = est),colour="grey70") +
  geom_errorbar(data=blackresp.rep.hispanic, aes(ymin = lb, ymax = ub), width = 0.2,
  colour="grey70") +
  geom_point(data= blackresp.rep.white, aes(x = model, y = est),colour="grey70") +
  geom_errorbar(data=blackresp.rep.white, aes(ymin = lb, ymax = ub), width = 0.2,
  colour="grey70")  +
  geom_point(data= whiteresp.rep.hispanic, aes(x = model, y = est),colour="black") +
  geom_errorbar(data=whiteresp.rep.hispanic, aes(ymin = lb, ymax = ub), width = 0.2,
  colour="black") +
  geom_point(data= whiteresp.rep.black, aes(x = model, y = est),colour="black") +
  geom_errorbar(data=whiteresp.rep.black, aes(ymin = lb, ymax = ub), width = 0.2,
  colour="black")  +
  theme_bw() + coord_flip()  +
  labs(title="Republicans: Support for Nominees by Race") +
  theme(plot.title = element_text(hjust = 0.5)) + ylim(-1.75,1.5)

grid.arrange(g3, g4, ncol=1, left= textGrob("Respondent-Candidate Pairing", rot=90),
             bottom=textGrob("Coefficient Estimate"))
```

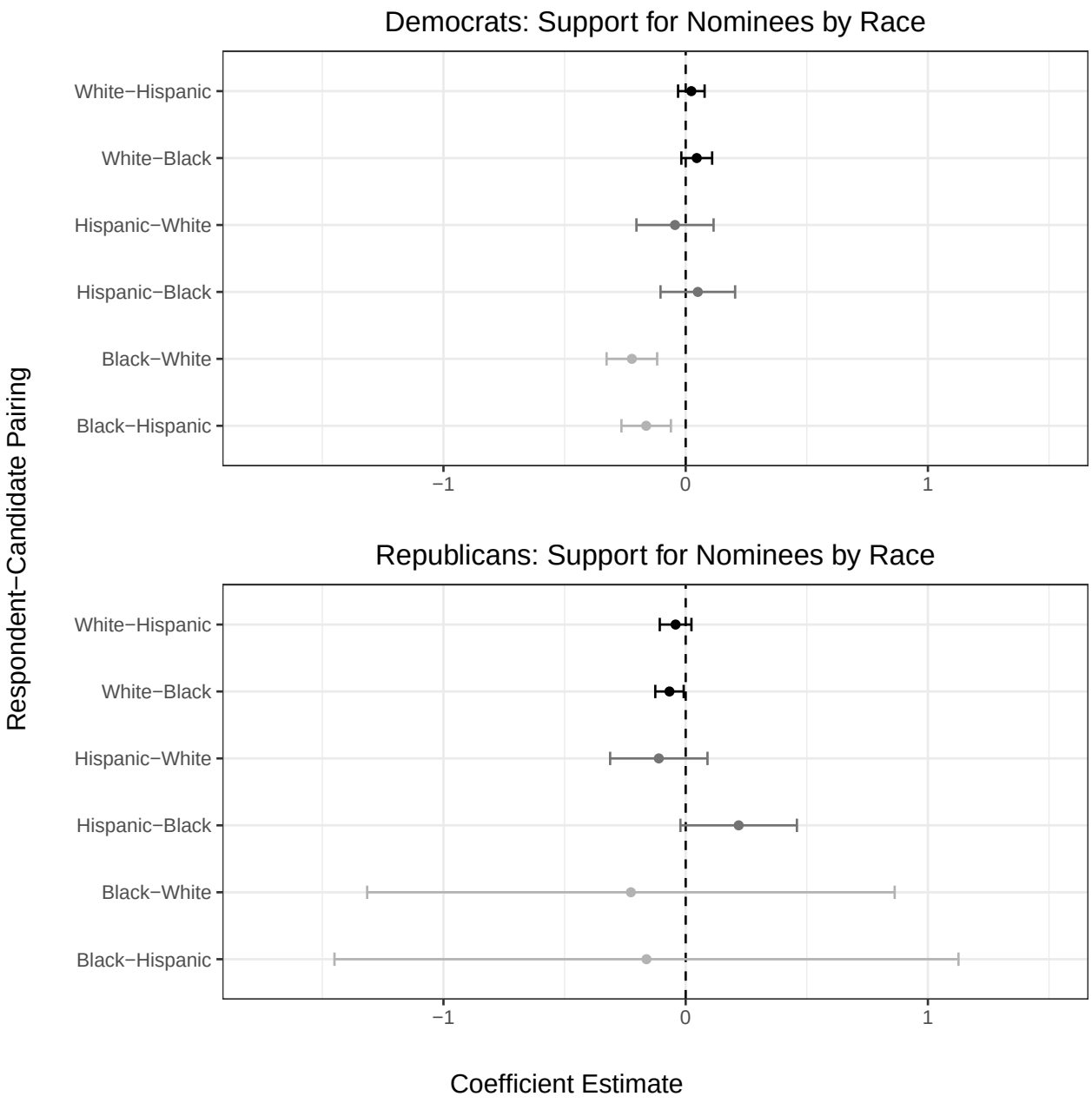
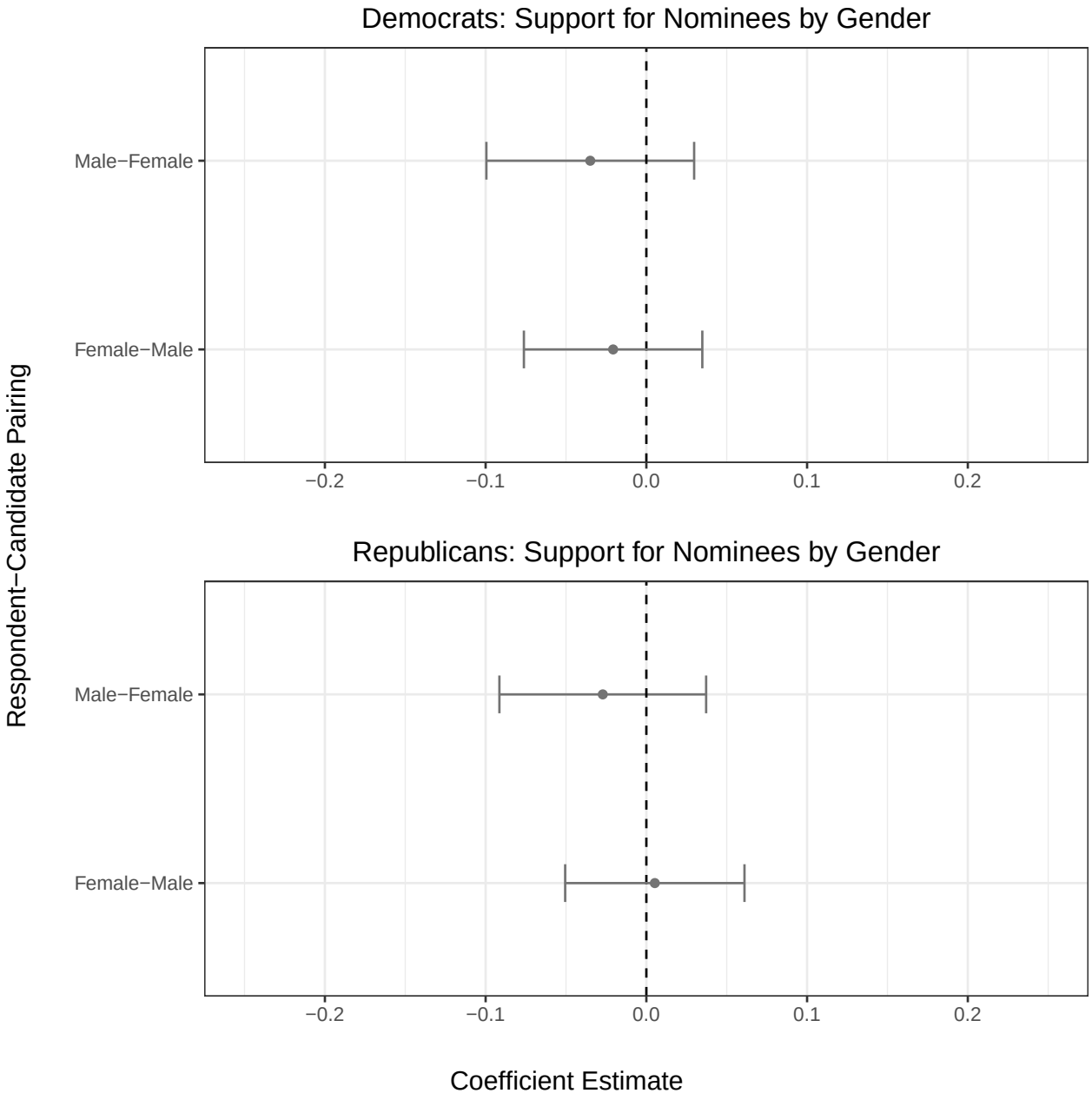


Figure C.3: Partisanship, Gender, and Nominee Support (Bootstrapped Standard Errors)

```
# Graph the coefficients and their confidence intervals
maleresp.dem.female <- data.frame(est = supportresults.mendems_vec_est["gendernominee"],
                                  lb =  supportresults.mendems_vec_est["gendernominee"] - qnorm(.975)*su
                                  ub = supportresults.mendems_vec_est["gendernominee"] + qnorm(.975)*sup
                                  model = "Male-Female")


femaleresp.dem.male <- data.frame(est = supportresults.womendems_vec_est["gendernominee"],
```

```
                                  lb =  supportresults.womendems_vec_est["gendernominee"] - qnorm(.975)*
                                  ub =  supportresults.womendems_vec_est["gendernominee"] + qnorm(.975)*
                                  model = "Female-Male")

maleresp.rep.female <- data.frame(est = supportresults.menreps_vec_est["gendernominee"],
                                  lb =  supportresults.menreps_vec_est["gendernominee"] - qnorm(.975)*su
                                  ub = supportresults.menreps_vec_est["gendernominee"] + qnorm(.975)*sup
                                  model = "Male-Female")


femaleresp.rep.male <- data.frame(est = supportresults.womenreps_vec_est["gendernominee"],
                                  lb = supportresults.womenreps_vec_est["gendernominee"] - qnorm(.975)*s
                                  ub = supportresults.womenreps_vec_est["gendernominee"] + qnorm(.975)*s
                                  model = "Female-Male")

g1 <- ggplot(maleresp.dem.female, aes(x = model, y = est)) +
  geom_point(colour="grey45") +
  geom_errorbar(aes(ymin = lb, ymax = ub), width = 0.2, colour="grey45") +
  geom_hline(yintercept = 0, lty = 2) +
  xlab("") +
  ylab("") +
  geom_point(data= femaleresp.dem.male, aes(x = model, y = est),
  colour="grey45") +
  geom_errorbar(data=femaleresp.dem.male, aes(ymin = lb, ymax = ub), width = 0.2,
  colour="grey45") +
  theme_bw() + coord_flip() + ylim(-.25,0.25) +
  labs(title="Democrats: Support for Nominees by Gender") +
  theme(plot.title = element_text(hjust = 0.5))


g2 <- ggplot(maleresp.rep.female, aes(x = model, y = est)) +
  geom_point(colour="grey45") +
  geom_errorbar(aes(ymin = lb, ymax = ub), width = 0.2,
  colour="grey45") +
  geom_hline(yintercept = 0, lty = 2) +
  xlab("") + ylab("") +
  geom_point(data= femaleresp.rep.male, aes(x = model, y = est),
  colour="grey45") +
  geom_errorbar(data=femaleresp.rep.male, aes(ymin = lb, ymax = ub),
  width = 0.2, colour="grey45") +
  theme_bw() + coord_flip() + ylim(-.25,0.25) +
  labs(title="Republicans: Support for Nominees by Gender") +
  theme(plot.title = element_text(hjust = 0.5))

grid.arrange(g1, g2, ncol=1, left= textGrob("Respondent-Candidate Pairing", rot=90),
             bottom=textGrob("Coefficient Estimate"))
```


Democrats: Support for Nominees by Gender
Male−Female
Female−Male
−0.2
−0.1
0.0
0.1
0.2
Republicans: Support for Nominees by Gender
Male−Female
Female−Male
−0.2
−0.1
0.0
0.1
0.2
Respondent−Candidate Pairing
Coefficient Estimate