```
fithigh11 <- glm( data = polbr2, formula = " high_interest ~ femrightsenough ", family ="binomial", weights = pesos, na.action = na.omit)

fithigh12 <- glm( data = polbr2, formula = " high_interest ~ parityagree ", family ="binomial", weights = pesos, na.action = na.omit)

fithigh13 <- glm( data = polbr2, formula = " high_interest ~ quotapunish ", family ="binomial", weights = pesos, na.action = na.omit)

fithigh14 <- glm( data = polbr2, formula = " high_interest ~ quotasenathird ", family ="binomial", weights = pesos, na.action = na.omit)

fithigh15 <- glm( data = polbr2, formula = " high_interest ~ haspartner ", family ="binomial", weights = pesos, na.action = na.omit)

fithigh16 <- glm( data = polbr2, formula = " high_interest ~ housekeep ", family ="binomial", weights = pesos, na.action = na.omit)

fithigh17 <- glm( data = polbr2, formula = " high_interest ~ haskids ", family ="binomial", weights = pesos, na.action = na.omit)

fithigh18 <- glm( data = polbr2, formula = " high_interest ~ haskidsin ", family ="binomial", weights = pesos, na.action = na.omit)

fithigh19 <- glm( data = polbr2, formula = " high_interest ~ haskidsinlittle ", family ="binomial", weights = pesos, na.action = na.omit)

fithigh20 <- glm( data = polbr2, formula = " high_interest ~ kidscare ", family ="binomial", weights = pesos, na.action = na.omit)

fithigh21 <- glm( data = polbr2, formula = " high_interest ~ kidstotal ", family ="binomial", weights = pesos, na.action = na.omit)

fithigh22 <- glm( data = polbr2, formula = " high_interest ~ wannakids ", family ="binomial", weights = pesos, na.action = na.omit)

fithigh23 <- glm( data = polbr2, formula = " high_interest ~ paidjob ", family ="binomial", weights = pesos, na.action = na.omit)

fithigh24 <- glm( data = polbr2, formula = " high_interest ~ assoc ", family ="binomial", weights = pesos, na.action = na.omit)

fithigh25 <- glm( data = polbr2, formula = " high_interest ~ levelassoc ", family ="binomial", weights = pesos, na.action = na.omit)

fithigh26 <- glm( data = polbr2, formula = " high_interest ~ familysupport ", family ="binomial", weights = pesos, na.action = na.omit)

fithigh27 <- glm( data = polbr2, formula = " high_interest ~ partnersupport ", family ="binomial", weights = pesos, na.action = na.omit)

fithigh28 <- glm( data = polbr2, formula = " high_interest ~ partnchangcitsupport ", family ="binomial", weights = pesos, na.action = na.omit)

fithigh29 <- glm( data = polbr2, formula = " high_interest ~ candwchance ", family ="binomial", weights = pesos, na.action = na.omit)

fithigh30 <- glm( data = polbr2, formula = " high_interest ~ thinkcand ", family ="binomial", weights = pesos, na.action = na.omit)

fithigh31 <- glm( data = polbr2, formula = " high_interest ~ sex ", family ="binomial", weights = pesos, na.action = na.omit)

fithigh32 <- glm( data = polbr2, formula = " high_interest ~ ageinterval ", family ="binomial", weights = pesos, na.action = na.omit)

fithigh33 <- glm( data = polbr2, formula = " high_interest ~ educalevel ", family ="binomial", weights = pesos, na.action = na.omit)

fithigh34 <- glm( data = polbr2, formula = " high_interest ~ incomeinterval ", family ="binomial", weights = pesos, na.action = na.omit)

fithigh35 <- glm( data = polbr2, formula = " high_interest ~ religion ", family ="binomial", weights = pesos, na.action = na.omit)

fithigh36 <- glm( data = polbr2, formula = " high_interest ~ tem_email ", family ="binomial", weights = pesos, na.action = na.omit)

fit1high_medsex <- glm( data = polbr2, formula = "high_med_interest ~ sex", family = "binomial", weights = pesos, na.action = na.omit)

summary(fit1high_medsex)

fit1medsex <- glm( data = polbr2, formula = "med_interest ~ sex", family = "binomial", weights = pesos, na.action = na.omit)

summary(fit1medsex)

fit1lowsex <- glm( data = polbr2, formula = "low_interest ~ sex", family = "binomial", weights = pesos, na.action = na.omit)

fit1lowtonosex <- glm( data = polbr2, formula = "low_no_interest ~ sex", family = "binomial", weights = pesos, na.action = na.omit)

fit1nosex <- glm( data = polbr2, formula = "sem_interest ~ sex", family = "binomial", weights = pesos, na.action = na.omit)

summary(fit1nosex)

summary(fit1nosex)

summary( fithigh1 ) summary( fithigh2 ) summary( fithigh3 ) summary( fithigh4 ) summary( fithigh5 ) summary( fithigh6 ) summary( fithigh7 ) summary( fithigh8 ) summary( fithigh9 ) summary( fithigh10 ) summary( fithigh11 ) summary( fithigh12 ) summary( fithigh13 ) summary( fithigh14 ) summary( fithigh15 ) summary( fithigh16 ) summary( fithigh17 ) summary( fithigh18 ) summary( fithigh19 ) summary( fithigh20 ) summary( fithigh21 ) summary( fithigh22 ) summary( fithigh23 ) summary( fithigh24 ) summary( fithigh25 ) summary( fithigh26 ) summary( fithigh27 ) summary( fithigh28 ) summary( fithigh29 ) summary( fithigh30 ) summary( fithigh31 ) summary( fithigh32 ) summary( fithigh33 ) summary( fithigh34 ) summary( fithigh35 ) summary( fithigh36 )
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Os resultados foram interessantes, mas parecem ter sido acometidos de erro de especificação. A suspeita decorre do fato de que simplismente todas as variáveis testadas se apresentaram estatisticamente significativas para explicar o alto interesse em política, quando pareadas de duas em duas.

Nenhum dos modelos com apenas uma variável explicativa (fithigh1 a fithigh36) apontou com clareza quais variáveis deveriam ir-se adicionando, nem permitiram fazer a seleção de variáveis que deveriam ser cortadas, com base no critério no p-valor.