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1  log using "22_analysis_log.txt", text replace
2  //// RCT Survey: Analysis
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4  //// 2021 April 24
5
6  cls
7  clear
8  set more off
9
10 cd "E:\lucas\projects_private_git\academia\projects\vaccine\rct"
11
12 use "data\data_shuffled_cleaned_2021-04-23_0939.dta", clear
13
14 **** Drop some variables that we're not using at this point in time
15 drop *_pagesubmit
16
17 *****
18 **** Generate variables
19 *****
20
21 **** Define segment and control variable lists
22 local segments "d_b d_l d_c d_r d_p"
23 local indivchars "d_male d_age_lt45 d_age_gt65 d_smalltown_rural"
24
25 **** Recommendation indicators
26 gen m_rec_b = ( m_rec_lebron | m_rec_kizzy | m_rec_obama | m_rec_warnock )
27 gen m_rec_l = ( m_rec_fernandez | m_rec_jlo | m_rec_badbunny )
28 gen m_rec_c = ( m_rec_trump )
29 gen m_rec_r = ( m_rec_warnock | m_rec_pope | m_rec_rickwarren )
30 gen m_rec_b_r = ( m_rec_b & m_rec_r ) // This is the only non-null intersection
31
32 **** Concordant recommendation
33 // Notice that here we subtract off m_rec_b_r if both d_b and d_r.
34 // That is, If a Black religious person gets a Warnock message, it obtains 1 point, not 2.
35 // Accordingly, concordant_rec is either 0 if not concordant or 1 if concordant.
36 gen concord_recommended = d_b*m_rec_b + d_l*m_rec_l + d_c*m_rec_c + d_r*m_rec_r - d_b*d_r*m_rec_b_r
37 tab concord_recommended
38
39 **** Concordant by message element
40 // Note that we need not multiply each message indicator by the demographic
41 // indicator because the messages were targeted to only those demographics.
42 // The one exception is m_frdm_c_r; this was shown to (d_c | d_r), but here we
43 // only count it as concordant if (d_c & d_r), so we use m_frdm_c_r_ANDdcANDdr.
44 gen concord_commimpact = m_comm_b + m_comm_l + m_comm_c + m_chrc_r
45 gen concord_poptested = m_popl_b + m_popl_l
46 gen concord_children = m_chld_p + m_chld_p_l
47 gen concord_elders = m_eldr_c + m_eldr_r + m_eldr_p
48 gen concord_protection = m_prtx_c
49 gen concord_gatherings = m_gthr_l + m_gthr_r + m_gthr_p + m_frdm_c_r_ANDdcANDdr
50 gen concord_availability = m_avyl_r + m_avyl_p
51
52 **** Count total concordant messages and recommendation
53 gen num_concord_msg = ///
54     concord_commimpact + concord_poptested + concord_children ///
55     + concord_elders + concord_protection + concord_gatherings ///
56     + concord_availability + concord_recommended
57
58 **** Create concordant score
59 // Give one point for each concordant message received.
60 // Give an extra point if Latinx and treated with Spanish parallel text.
61 // (Note that only Latinx receives Spanish, so no need to multiply by d_l.)
62 gen concordant_score = num_concord_msg + t_spanish
63
64 **** Concordant messages by segment
65 ** Black segment: community impact, population tested, recommendation
66 gen concord_b = m_comm_b + m_popl_b + d_b*m_rec_b
67 ** Latinx segment: comm. impact, pop. tested, gathering, recommendation
68 // We do not include m_chld_p_l because it was only a modification of m_chld_p
69 // for Latinx; the Latinx segment did not obtain a control of m_chld_p.
70 gen concord_l = m_comm_l + m_popl_l + m_gthr_l + d_l*m_rec_l
71 ** Conservative segment: comm. impact, elders, protection, freedom
72 // Again here we only count the m_frdm_c_r message as concordant if the subject
73 // is both conservative and religious.
74 gen concord_c = m_comm_c + m_eldr_c + m_prtx_c + m_frdm_c_r_ANDdcANDdr
75 ** Religious segment: comm. impact (church), elders, gatherings, avail., rec.
76 // To avoid double-counting m_frdm_c_r, we omit it from the religious segment
77 // and only include it in the conservative segment above.
78 gen concord_r = m_chrc_r + m_eldr_r + m_gthr_r + m_avyl_r + d_r*m_rec_r

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79  ** Parent segment: children, elders, gatherings, availability
80  // We can simply use (m_chld_p + m_chld_p_1) because no one received both.
81  gen concord_p = m_chld_p + m_chld_p_1 + m_eldr_p + m_gthr_p + m_avyl_p
82
83  *****
84  **** Drop useless observations
85  *****
86
87  // drop outcomes of "Don't know / Prefer not to say"
88  drop if vax_likely==8
89
90  *****
91  **** Drop subsegments with 10 or fewer obs.
92  *****
93
94  tab subseg
95
96  bysort subseg: gen subseg_N = _N
97  drop if subseg_N <= 10
98
99  tab subseg
100
101  *****
102  *****
103  **** ANALYSIS
104  *****
105  *****
106
107  *****
108  **** Descriptive
109  *****
110
111  **** Descriptive tables
112  // We'll produce many more tables than those here
113  tab d_relig d_r
114
115  **** Descriptive regressions on segment
116  ologit vax_likely `segments'
117  ologit vax_likely `segments' `indivchars'
118
119  **** Descriptive regressions on segment
120  ologit vax_likely i.subseg
121  ologit vax_likely i.subseg `indivchars'
122
123  *****
124  **** Primary analysis
125  *****
126
127  **** Likelihood to vaccinate = F(concordant_score)
128
129  reg vax_likely concordant_score i.subseg, cluster(subseg)
130  eststo e1
131
132  ologit vax_likely concordant_score i.subseg, cluster(subseg)
133  // Average marginal effects using margins
134  margins, dydx(concordant_score)
135  eststo e2
136
137  esttab e1 e2 using "30_results.txt", ci label addnote("Likelihood 1 to 7") append ///
138  title("How does adding an additional concordant message affect likelihood to vaccinate?")
139  eststo clear
140
141  ** Typical change in Pr(each response) as concordant_score rises
142  // https://www3.nd.edu/~rwilliam/xsoc73994/Margins05.pdf
143  *findit spost13_ado
144
145  ** Shows the effect of one more concordant_score point on Pr(each value of vax_likely)
146  // Due to a bug, we have to remove the variable's label (we add it back afterward)
147  label values vax_likely .
148  mtable, dydx(concordant_score)
149  label values vax_likely vax_likely_label
150
151  **** Likelihood to vaccinate = F(concordant score by topic)
152  reg vax_likely concord_commimpact concord_poptested concord_children ///
153  concord_elders concord_protection concord_gatherings ///
154  concord_availability concord_recommended i.subseg, cluster(subseg)
155  est sto e1
156

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157 **** Likelihood to vaccinate = F(concordant score by segment)
158 reg vax_likely concord_b concord_l concord_c concord_r concord_p i.subseg, cluster(subseg)
159 est sto e2
160
161 esttab e1 e2 using "30_results.txt", ci label addnote("Likelihood 1 to 7") append ///
162     title("Which concordant topics matter? Which subsegments respond to concordance?")
163 eststo clear
164
165 *****
166 **** Close log
167 *****
168
169 log close
170
```