Replication 3

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R Markdown

Adding additional functions

Reading in the data

Cleaning up the data

Loading in some formulas

Figure 2

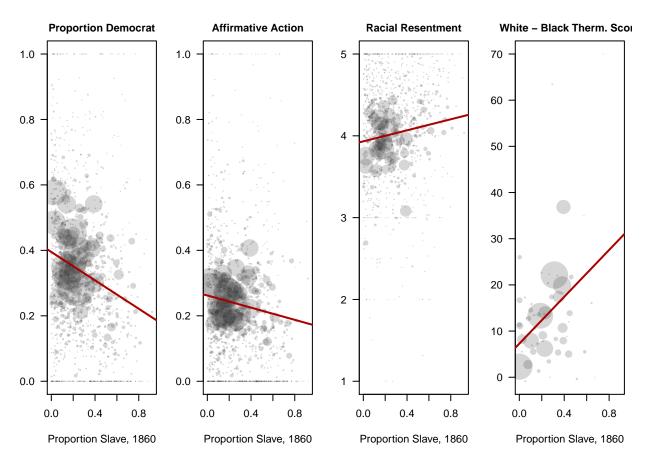


Table 1

	Prop Democrat		Affirm.		
	(1)	(2)	(3)	(4)	(5)
Prop. Slave, 1860	-0.217^{***} (0.023)	-0.127^{***} (0.043)	-0.145^{***} (0.036)	0.463*** (0.139)	36.125*** (9.244)
Level 1860 Covariates State Fixed Effects State-Year Fixed Effects Clustered SEs	County	County ✓	County ✓	County ✓	Individual ✓ ✓ ✓
$rac{N}{R^2}$	$1,242 \\ 0.065$	$1,152 \\ 0.203$	$1,152 \\ 0.101$	$1,027 \\ 0.143$	$1,489 \\ 0.183$

p < .1; p < .05; p < .01

Note. Models 1–4 are weighted least squares at the county level, with within-county sample sizes, adjusted by sampling weights, as weights. Model 5 is weighted least squares at the individual level with state-year fixed effects. Standard errors, in parentheses, are clustered at the county level in model 5. There are 50 counties in model 5.

 ${\bf Table~2}$ The data contain missing values. CEM will match on them; see the manual for other options.

	Prop Democrat						
	(1)	(2)	(3)	(4)	(5)	(6)	
Prop. Slave, 1860	-0.202^{***} (0.068)		-0.149^{***} (0.056)		0.450** (0.228)		
Slave State	,	$0.078* \\ (0.046)$,	-0.091^{**} (0.041)	,	-0.047 (0.149)	
State Fixed Effects	\checkmark		\checkmark		\checkmark		
1860 Covariates	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	
50% Threshold Match North-South Match	✓	\checkmark	✓	\checkmark	✓	\checkmark	
N	444	145	444	145	393	127	
\mathbb{R}^2	0.248	0.202	0.156	0.208	0.195	0.113	

p < .1; p < .05; p < .05; ***p < .01

Table 3

	Prop Democrat							
	(1)	(2)	(3)	(4)	(5)	(6)		
Prop. Slave, Direct Effect	-0.150^{***} (0.043)	-0.130^{**} (0.057)	-0.153^{***} (0.037)	-0.130^{***} (0.042)	0.514*** (0.140)	0.472** (0.184)		
Prop. Black, 2000	0.172^{***} (0.043)	(0.001)	0.062 (0.037)	(0.042)	$ \begin{array}{c} (0.140) \\ -0.416^{***} \\ (0.140) \end{array} $	(0.104)		
State Fixed Effects	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	✓		
1860 Covariates	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark	\checkmark		
Bootstrapped SEs		\checkmark		\checkmark		\checkmark		
Model	WLS	Seq. g-est.	WLS	Seq. g-est.	WLS	Seq. g-est.		
N	1,152	1,152	1,152	1,152	1,027	1,027		
\mathbb{R}^2	0.214	0.209	0.103	0.095	0.151	0.144		

p < .1; p < .05; p < .01