

**Catastrophic Success: Why Foreign-Imposed Regime Change Goes Wrong
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**Supplementary Materials for Chapter 3, Part 2: Genetic Matching¹
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¹ For information on the procedures used to create the matched datasets and perform the subsequent analyses, see pp. 150-51 of the book.

FIRC

Note: **peaceyrs** is excluded from matching because the balance between treated (33.48) and control (34.50) cases was already very good.

summary(firc.out2)

Call:

```
matchit(formula = abdfirctarg ~ lnpec4_fill + lntpop4_banks +
  mountainous + newstate + democracy + losewar10 + cw1000ongoing201411 +
  region1 + region2 + region3 + region4 + region5, data = firc,
  method = "genetic", pop.size = 150)
```

Summary of balance for all data:

	Means Treated	Means Control	SD Control	Mean Diff	eQQ Med	eQQ Mean	eQQ Max
distance	0.0991	0.0120	0.0286	0.0871	0.0336	0.0855	0.3584
lnpec4_fill	4.7369	7.0217	3.8684	-2.2848	1.8751	2.2934	5.7859
lntpop4_banks	8.0128	8.5581	1.8705	-0.5453	0.6369	0.6864	2.3136
mountainous	0.2299	0.1832	0.2210	0.0467	0.0635	0.0663	0.2135
newstate	0.0450	0.0256	0.1580	0.0194	0.0000	0.0180	1.0000
democracy	0.1441	0.3557	0.4788	-0.2116	0.0000	0.2162	1.0000
losewar10	0.4144	0.0518	0.2217	0.3626	0.0000	0.3604	1.0000
cw1000ongoing201411	0.1532	0.0908	0.2873	0.0624	0.0000	0.0631	1.0000
region1	0.4144	0.2224	0.4159	0.1920	0.0000	0.1892	1.0000
region2	0.0360	0.1566	0.3635	-0.1206	0.0000	0.1261	1.0000
region3	0.1261	0.2046	0.4034	-0.0785	0.0000	0.0811	1.0000
region4	0.1171	0.1910	0.3931	-0.0739	0.0000	0.0811	1.0000
region5	0.3063	0.2253	0.4178	0.0810	0.0000	0.0811	1.0000

Summary of balance for matched data:

	Means Treated	Means Control	SD Control	Mean Diff	eQQ Med	eQQ Mean	eQQ Max
distance	0.0991	0.0925	0.1083	0.0066	0.0114	0.0245	0.2528
lnpec4_fill	4.7369	5.1696	3.9045	-0.4327	0.2614	1.0107	4.5218
lntpop4_banks	8.0128	8.2304	1.3015	-0.2177	0.6119	0.6671	1.7792
mountainous	0.2299	0.2191	0.1894	0.0108	0.0139	0.0234	0.1175
newstate	0.0450	0.0450	0.2089	0.0000	0.0000	0.0139	1.0000
democracy	0.1441	0.1261	0.3343	0.0180	0.0000	0.0278	1.0000
losewar10	0.4144	0.4144	0.4961	0.0000	0.0000	0.0694	1.0000
cw1000ongoing201411	0.1532	0.1802	0.3870	-0.0270	0.0000	0.0694	1.0000
region1	0.4144	0.4414	0.5000	-0.0270	0.0000	0.0417	1.0000
region2	0.0360	0.0360	0.1877	0.0000	0.0000	0.0139	1.0000
region3	0.1261	0.0901	0.2883	0.0360	0.0000	0.0139	1.0000
region4	0.1171	0.1171	0.3238	0.0000	0.0000	0.0556	1.0000
region5	0.3063	0.3153	0.4679	-0.0090	0.0000	0.0278	1.0000

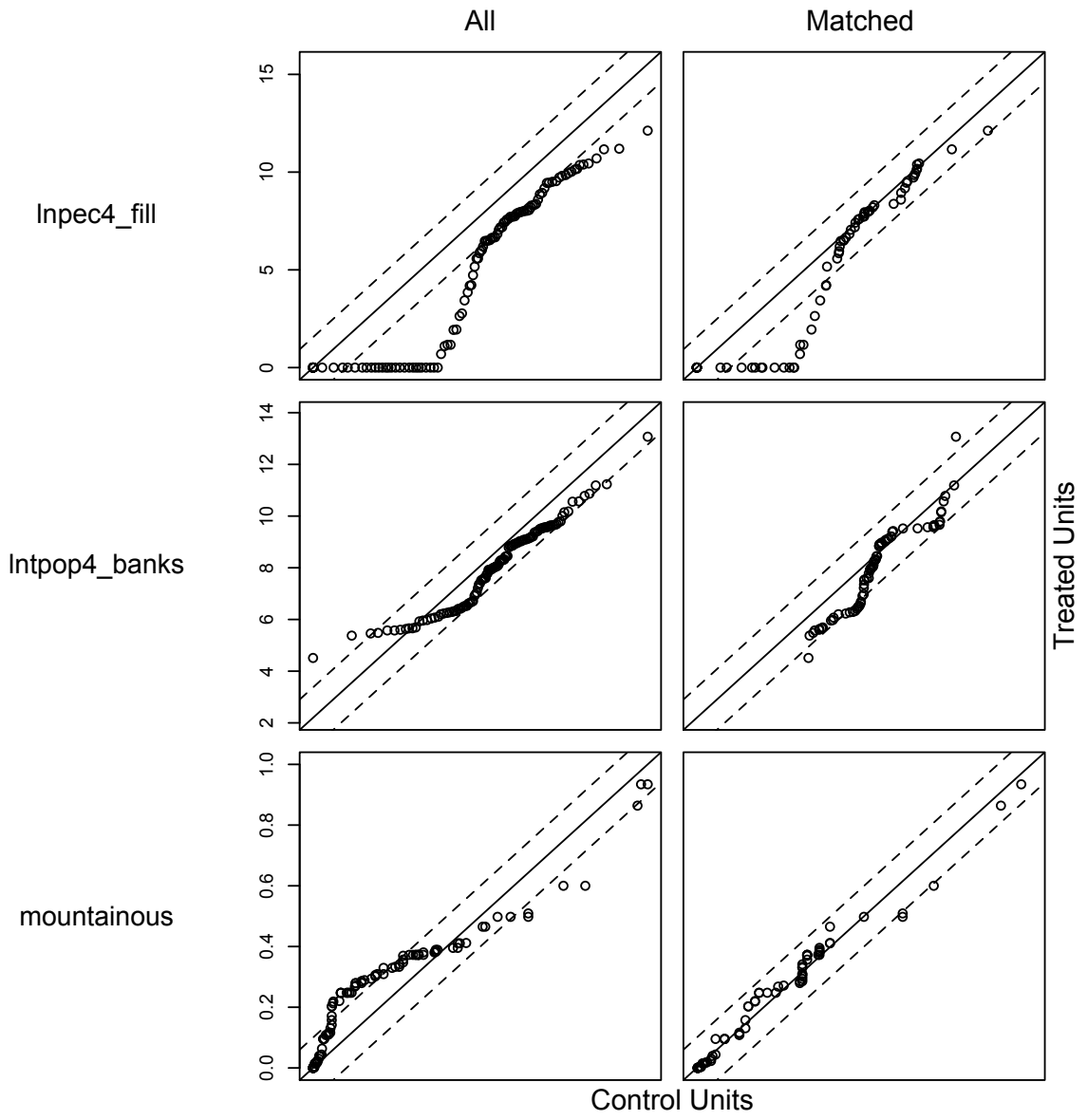
Percent Balance Improvement:

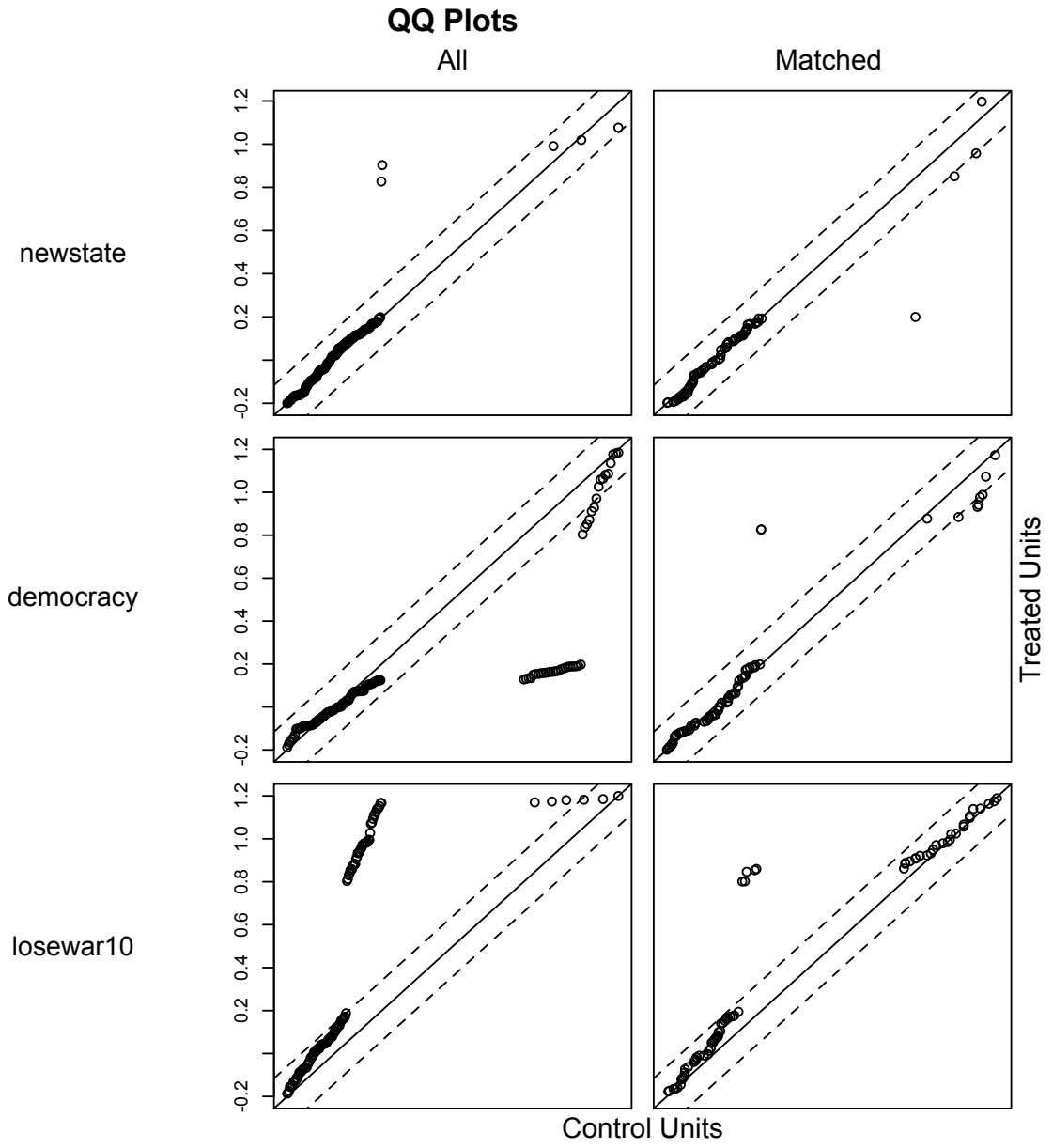
	Mean Diff.	eQQ Med	eQQ Mean	eQQ Max
distance	92.3728	66.0653	71.4134	29.4532
lnpec4_fill	81.0613	86.0620	55.9303	21.8480
lntpop4_banks	60.0846	3.9251	2.8091	23.0989
mountainous	76.8794	78.1911	64.6260	44.9670
newstate	100.0000	0.0000	22.9167	0.0000
democracy	91.4845	0.0000	87.1528	0.0000
losewar10	100.0000	0.0000	80.7292	0.0000
cw1000ongoing201411	56.6778	0.0000	-10.1190	0.0000
region1	85.9238	0.0000	77.9762	0.0000
region2	100.0000	0.0000	88.9881	0.0000
region3	54.0886	0.0000	82.8704	0.0000
region4	100.0000	0.0000	31.4815	0.0000
region5	88.8794	0.0000	65.7407	0.0000

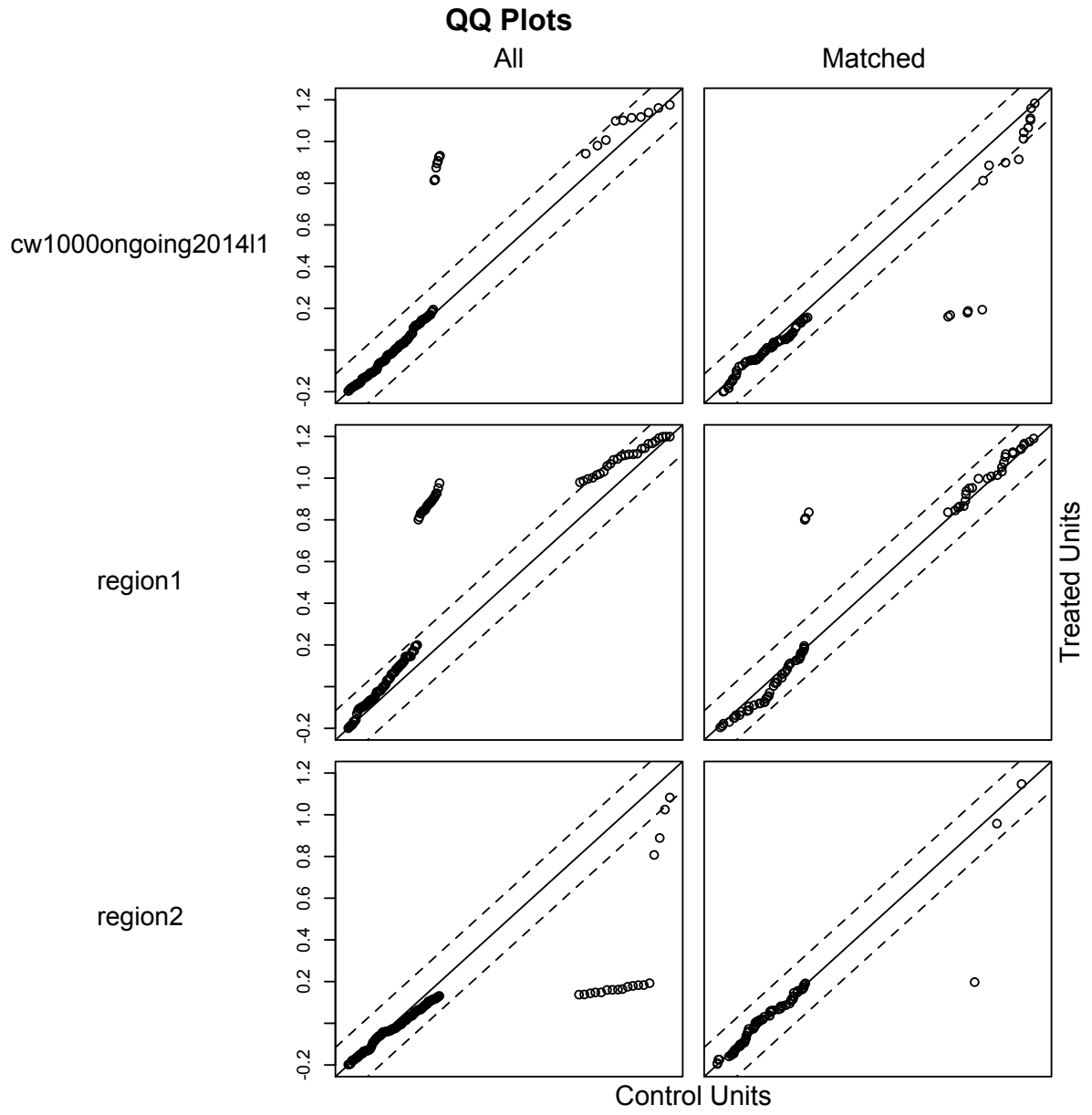
Sample sizes:

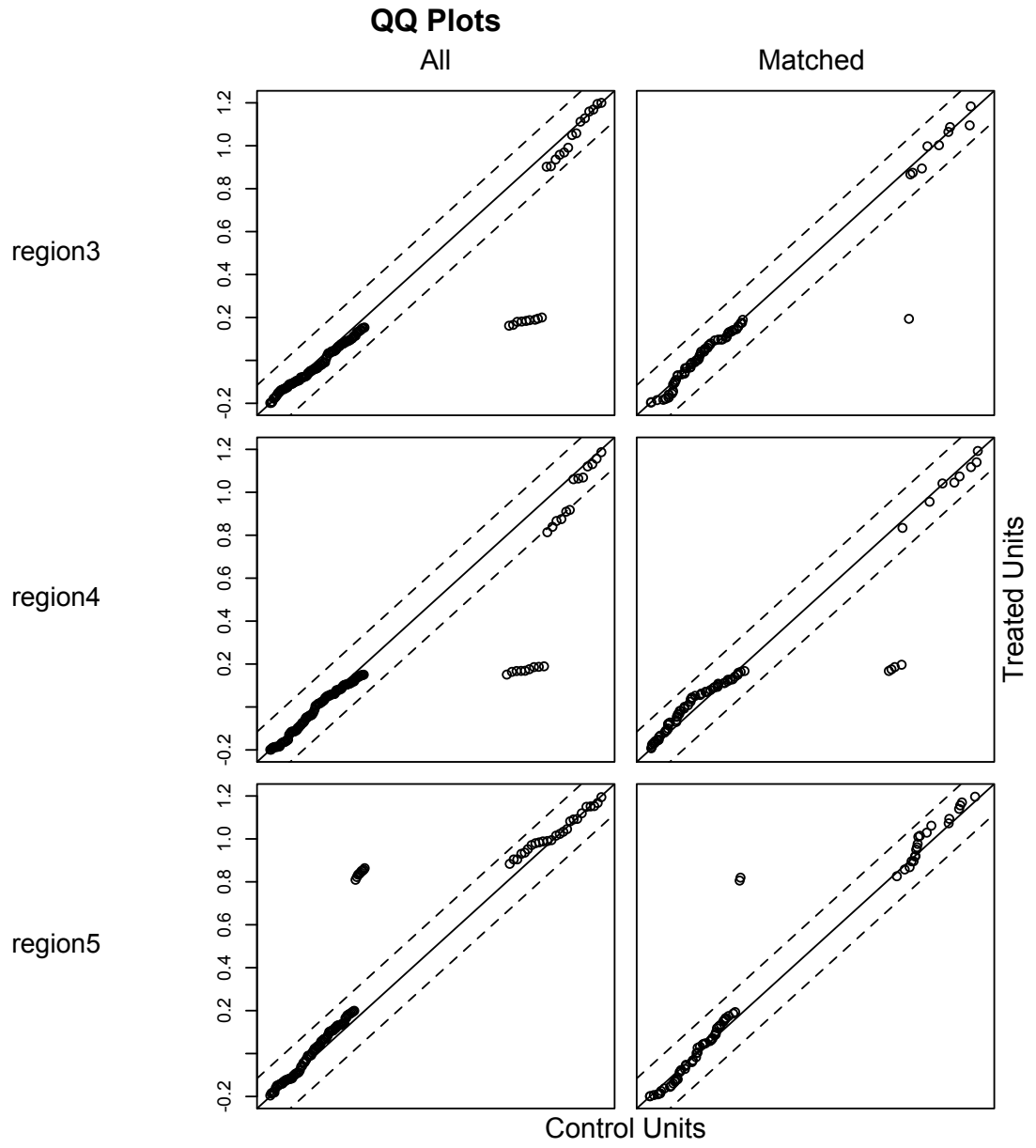
	Control	Treated
All	8318	111
Matched	72	111
Unmatched	8246	0
Discarded	0	0

QQ Plots

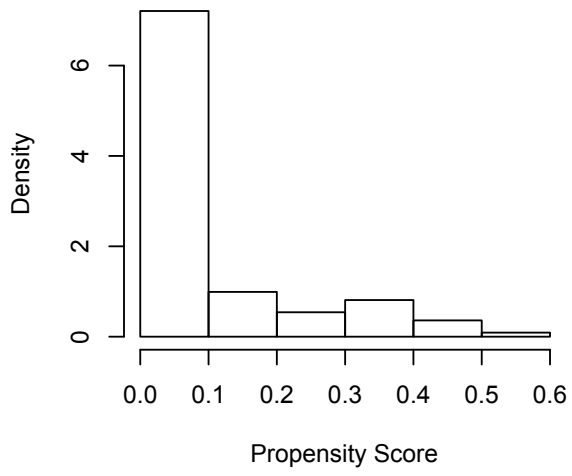




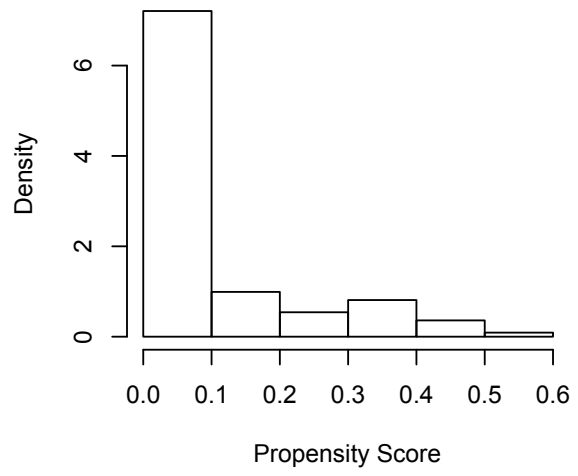




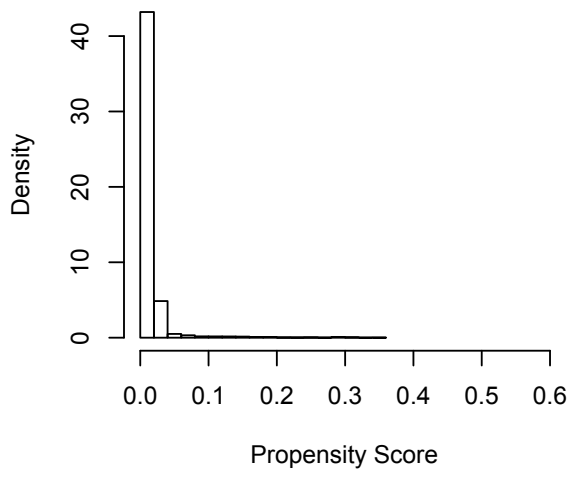
Raw Treated



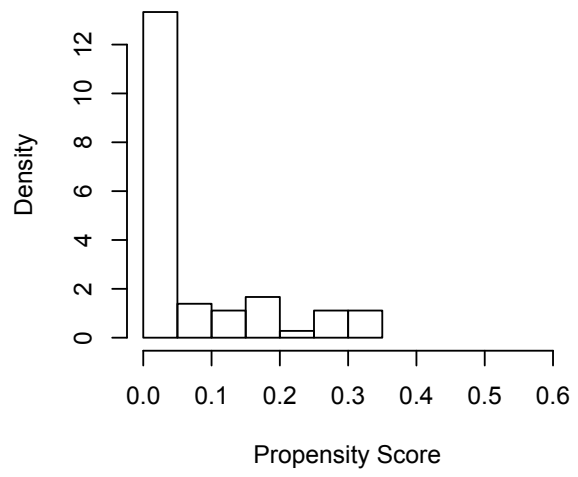
Matched Treated



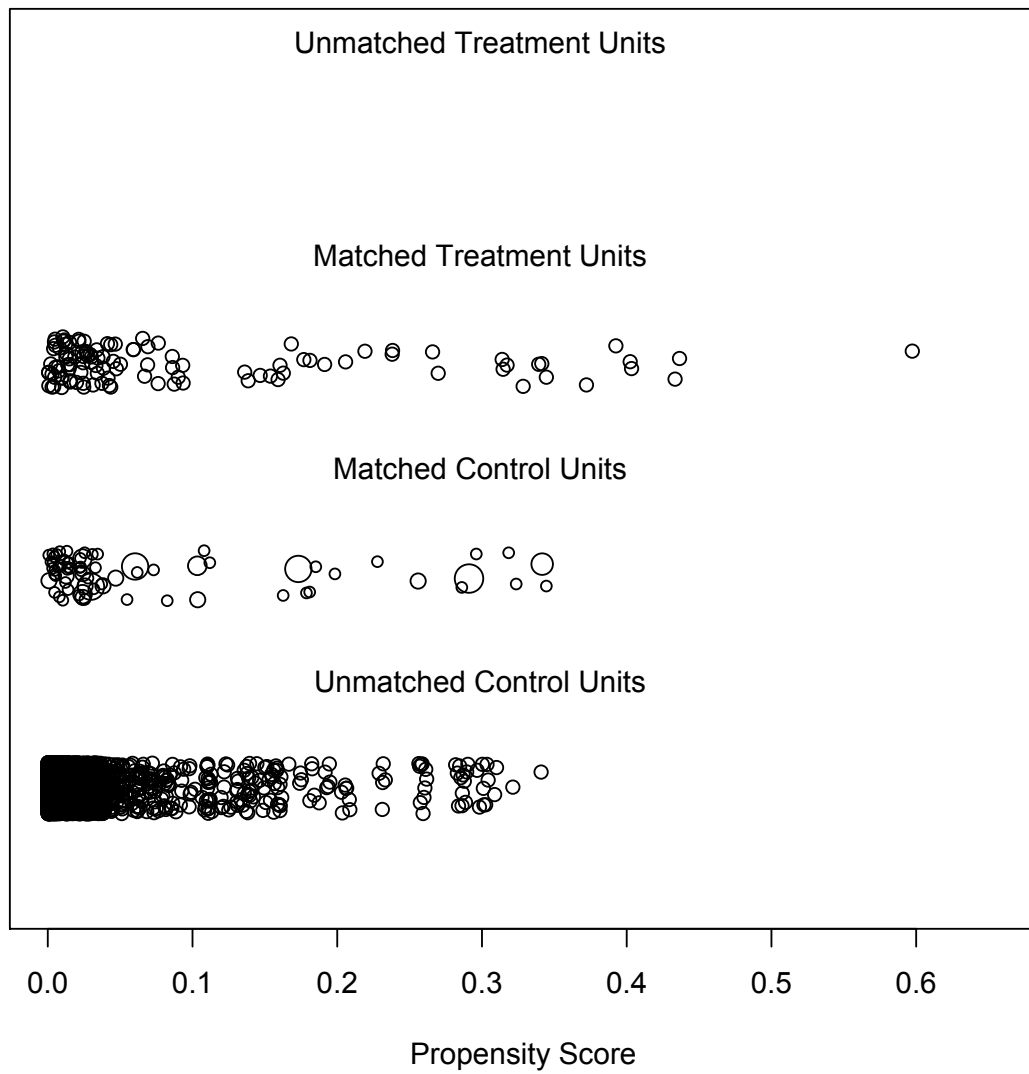
Raw Control



Matched Control



Distribution of Propensity Scores



FIRC: MATCHED ANALYSIS

tab civwar10_2014 abdfirtarg, col chi2

civwar10_2014	abdfirtarg		Total
	0	1	
0	61 84.72	75 67.57	136 74.32
1	11 15.28	36 32.43	47 25.68
Total	72 100.00	111 100.00	183 100.00

Pearson chi2(1) = 6.7334 Pr = 0.009

tab civwar5_2014 abdfirtarg, col chi2

civwar5_2014	abdfirtarg		Total
	0	1	
0	65 90.28	81 72.97	146 79.78
1	7 9.72	30 27.03	37 20.22
Total	72 100.00	111 100.00	183 100.00

Pearson chi2(1) = 8.1075 Pr = 0.004

ttest civwar10_2014, by(abdfirtarg)

Two-sample t test with equal variances

Group	Obs	Mean	Std. Err.	Std. Dev.	[95% Conf. Interval]	
0	72	.1527778	.0426972	.3622979	.0676419	.2379136
1	111	.3243243	.0446337	.4702449	.2358709	.4127778
combined	183	.2568306	.0323841	.4380838	.1929341	.3207271
diff		-.1715465	.0652395		-.3002744	-.0428187
diff = mean(0) - mean(1)					t =	-2.6295
Ho: diff = 0					degrees of freedom =	181

Ha: diff < 0 Ha: diff != 0 Ha: diff > 0
 Pr(T < t) = 0.0046 Pr(|T| > |t|) = 0.0093 Pr(T > t) = 0.9954

ttest civwar5_2014, by(abdfirtarg)

Two-sample t test with equal variances

Group	Obs	Mean	Std. Err.	Std. Dev.	[95% Conf. Interval]	
0	72	.0972222	.0351596	.2983392	.027116	.1673285
1	111	.2702703	.0423432	.4461134	.186356	.3541846
combined	183	.2021858	.0297708	.4027319	.1434455	.2609261
diff		-.173048	.0597407		-.2909258	-.0551703
diff = mean(0) - mean(1)					t =	-2.8967
Ho: diff = 0					degrees of freedom =	181

Ha: diff < 0 Ha: diff != 0 Ha: diff > 0
 Pr(T < t) = 0.0021 Pr(|T| > |t|) = 0.0042 Pr(T > t) = 0.9979

FIRC (using interwar and gdbuffer)

summary(firc.out3)

Call:

```
matchit(formula = abdfirctarg ~ lnpec4_fill + lnpop4_banks +
  mountainous + newstate + democracy + interwar + cw1000ongoing201411 +
  gdbuffer + region1 + region2 + region3 + region4 + region5,
  data = firc, method = "genetic", pop.size = 150)
```

Summary of balance for all data:

	Means	Treated	Means	Control	SD	Control	Mean Diff	eQQ	Med	eQQ	Mean	eQQ	Max
distance		0.0791		0.0123		0.0293	0.0668	0.0312		0.0650		0.2947	
lnpec4_fill		4.7369		7.0217		3.8684	-2.2848	1.8751		2.2934		5.7859	
lnpop4_banks		8.0128		8.5581		1.8705	-0.5453	0.6369		0.6864		2.3136	
mountainous		0.2299		0.1832		0.2210	0.0467	0.0635		0.0663		0.2135	
newstate		0.0450		0.0256		0.1580	0.0194	0.0000		0.0180		1.0000	
democracy		0.1441		0.3557		0.4788	-0.2116	0.0000		0.2162		1.0000	
interwar		0.3604		0.0513		0.2207	0.3090	0.0000		0.3063		1.0000	
cw1000ongoing201411		0.1532		0.0908		0.2873	0.0624	0.0000		0.0631		1.0000	
gdbuffer		0.3243		0.1193		0.3241	0.2051	0.0000		0.1982		1.0000	
region1		0.4144		0.2224		0.4159	0.1920	0.0000		0.1892		1.0000	
region2		0.0360		0.1566		0.3635	-0.1206	0.0000		0.1261		1.0000	
region3		0.1261		0.2046		0.4034	-0.0785	0.0000		0.0811		1.0000	
region4		0.1171		0.1910		0.3931	-0.0739	0.0000		0.0811		1.0000	
region5		0.3063		0.2253		0.4178	0.0810	0.0000		0.0811		1.0000	

Summary of balance for matched data:

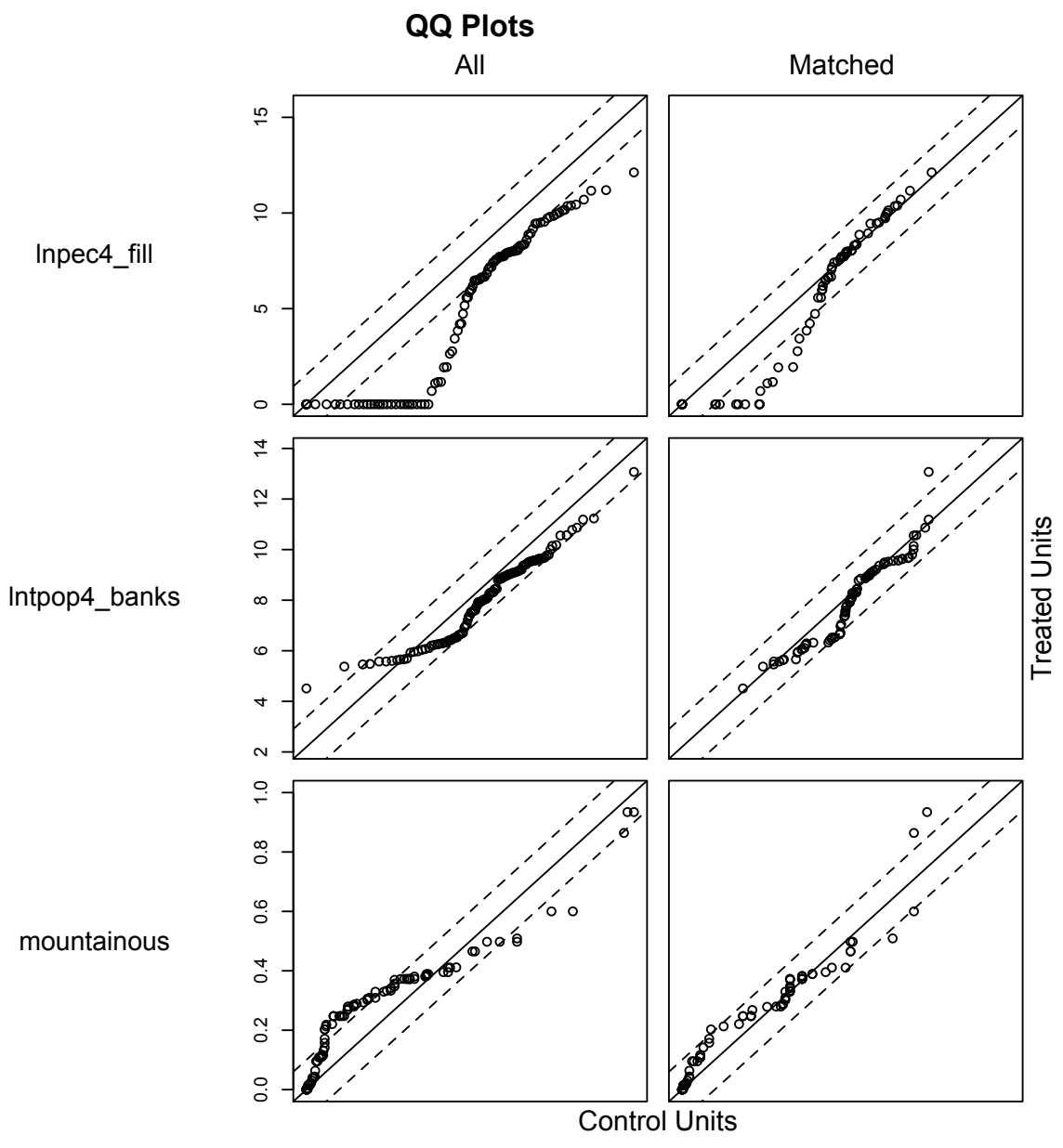
	Means	Treated	Means	Control	SD	Control	Mean Diff	eQQ	Med	eQQ	Mean	eQQ	Max
distance		0.0791		0.0748		0.0901	0.0043	0.0038		0.0085		0.0537	
lnpec4_fill		4.7369		4.7874		3.6376	-0.0506	0.2017		0.7675		3.6889	
lnpop4_banks		8.0128		8.0524		1.4586	-0.0396	0.2891		0.3893		2.0285	
mountainous		0.2299		0.2099		0.1968	0.0200	0.0173		0.0329		0.1862	
newstate		0.0450		0.0450		0.2087	0.0000	0.0000		0.0253		1.0000	
democracy		0.1441		0.1351		0.3441	0.0090	0.0000		0.0127		1.0000	
interwar		0.3604		0.3604		0.4832	0.0000	0.0000		0.0000		0.0000	
cw1000ongoing201411		0.1532		0.1532		0.3624	0.0000	0.0000		0.0253		1.0000	
gdbuffer		0.3243		0.3243		0.4711	0.0000	0.0000		0.0000		0.0000	
region1		0.4144		0.4144		0.4958	0.0000	0.0000		0.0127		1.0000	
region2		0.0360		0.0360		0.1876	0.0000	0.0000		0.0000		0.0000	
region3		0.1261		0.1261		0.3341	0.0000	0.0000		0.0380		1.0000	
region4		0.1171		0.1171		0.3236	0.0000	0.0000		0.0127		1.0000	
region5		0.3063		0.3063		0.4639	0.0000	0.0000		0.0253		1.0000	

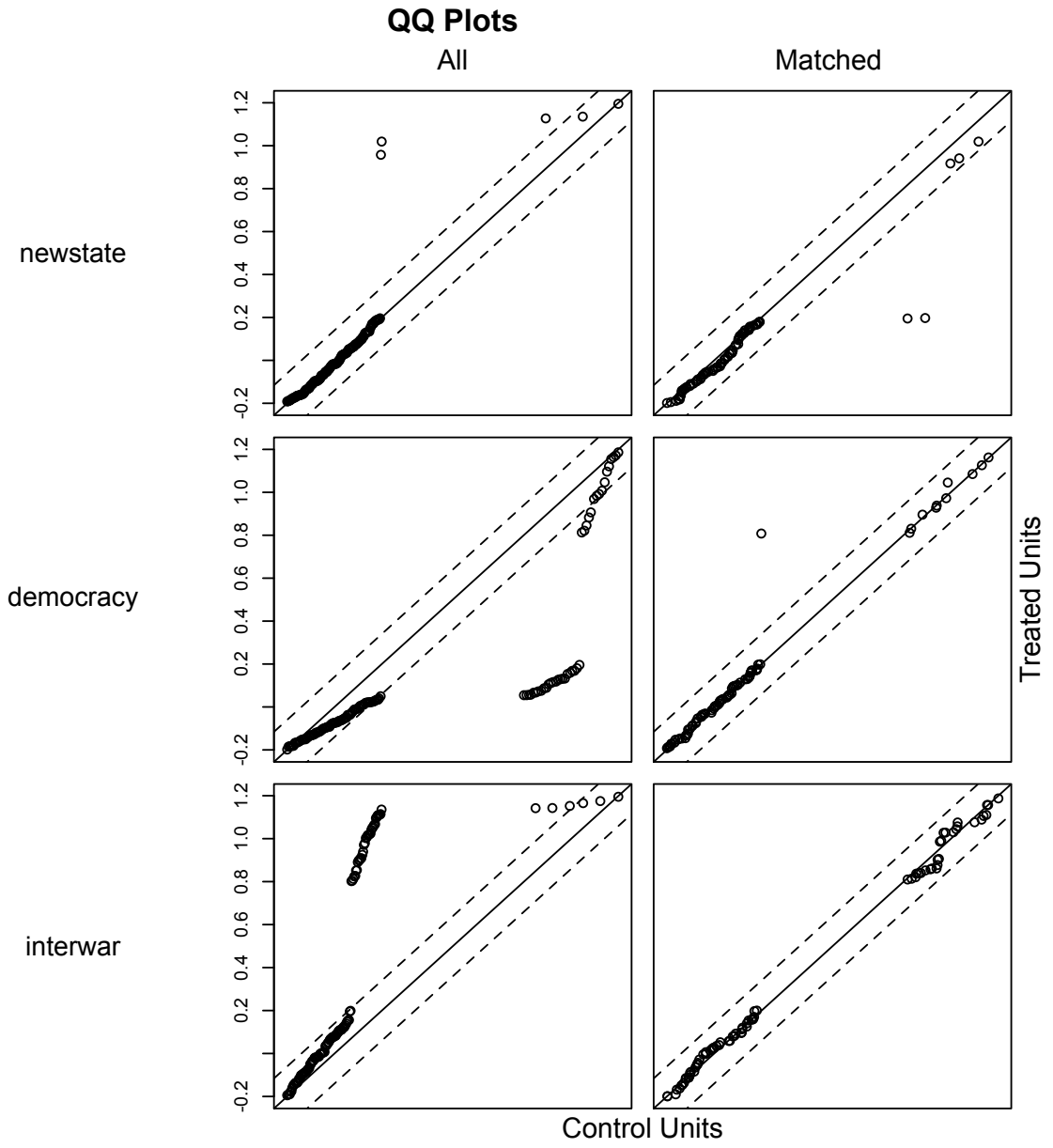
Percent Balance Improvement:

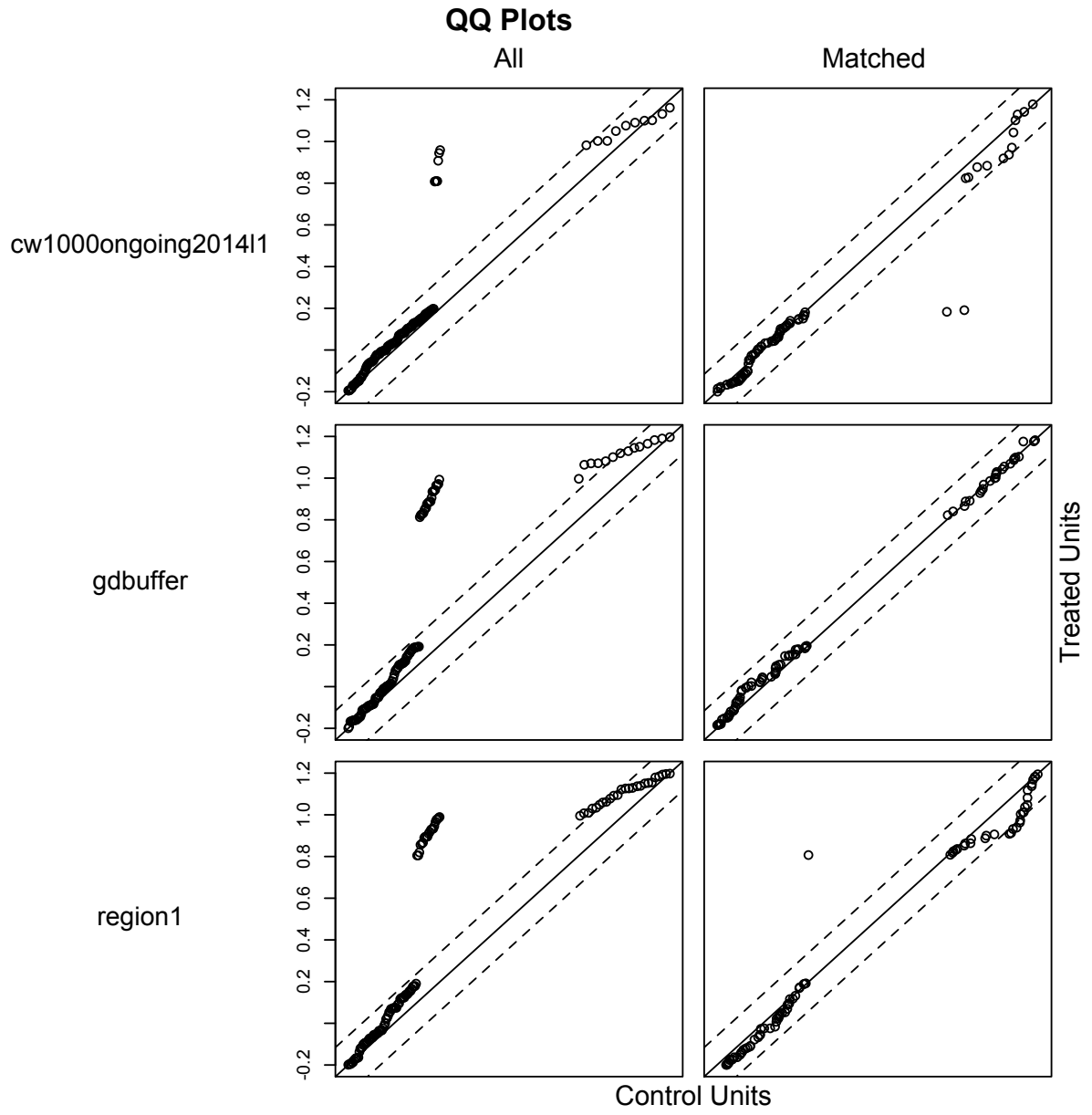
	Mean Diff.	eQQ	Med	eQQ	Mean	eQQ	Max
distance	93.5313	87.9372	86.8650	81.7699			
lnpec4_fill	97.7864	89.2451	66.5343	36.2435			
lnpop4_banks	92.7384	54.6068	43.2851	12.3254			
mountainous	57.2228	72.8214	50.4117	12.7931			
newstate	100.0000	0.0000	-40.5063	0.0000			
democracy	95.7422	0.0000	94.1456	0.0000			
interwar	100.0000	0.0000	100.0000	100.0000			
cw1000ongoing201411	100.0000	0.0000	59.8553	0.0000			
gdbuffer	100.0000	0.0000	100.0000	100.0000			
region1	100.0000	0.0000	93.3092	0.0000			
region2	100.0000	0.0000	100.0000	100.0000			
region3	100.0000	0.0000	53.1646	0.0000			
region4	100.0000	0.0000	84.3882	0.0000			
region5	100.0000	0.0000	68.7764	0.0000			

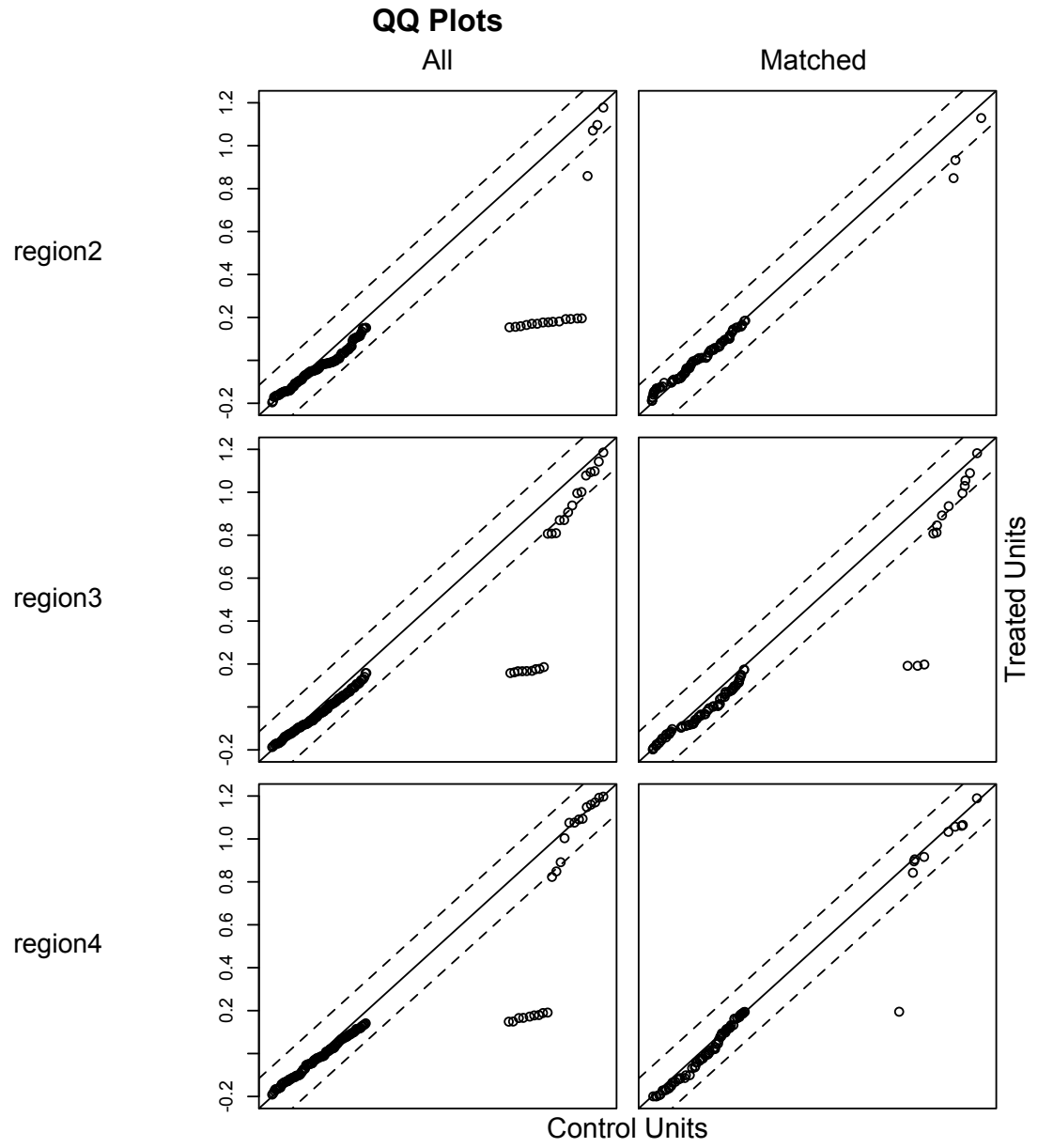
Sample sizes:

	Control	Treated
All	8318	111
Matched	79	111
Unmatched	8239	0
Discarded	0	0

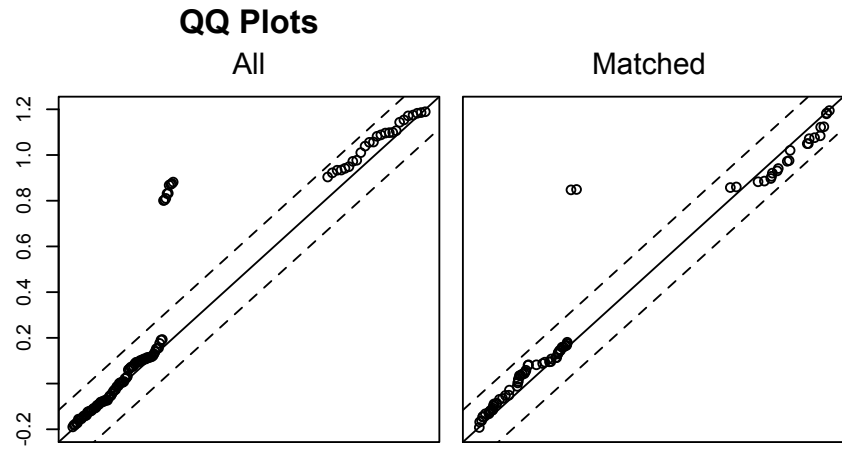








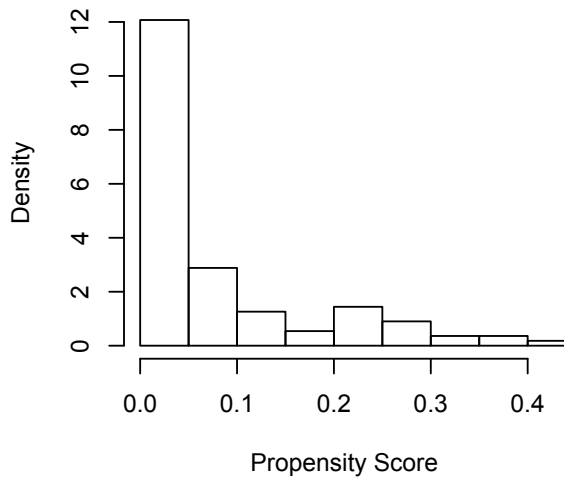
region5



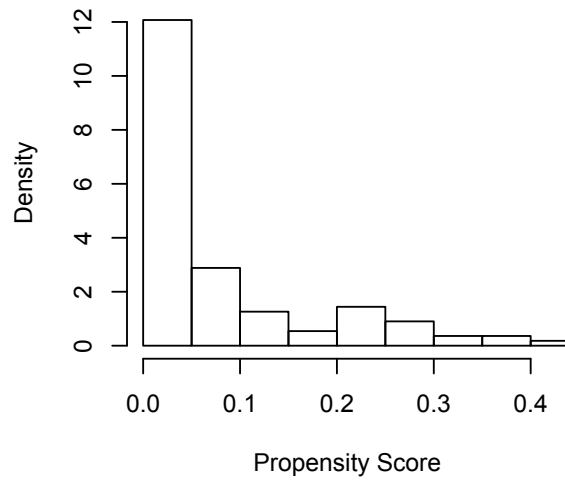
Treated Units

Control Units

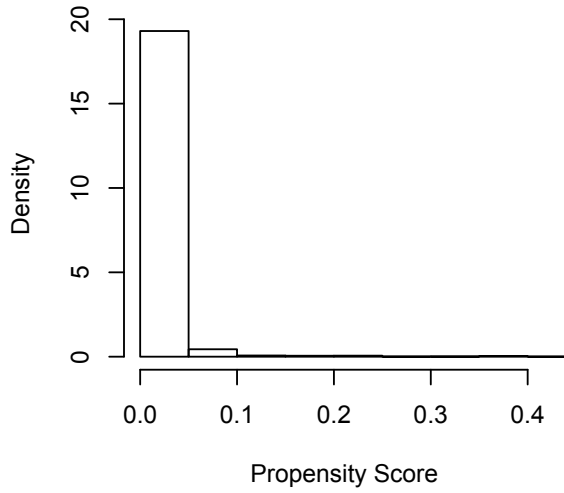
Raw Treated



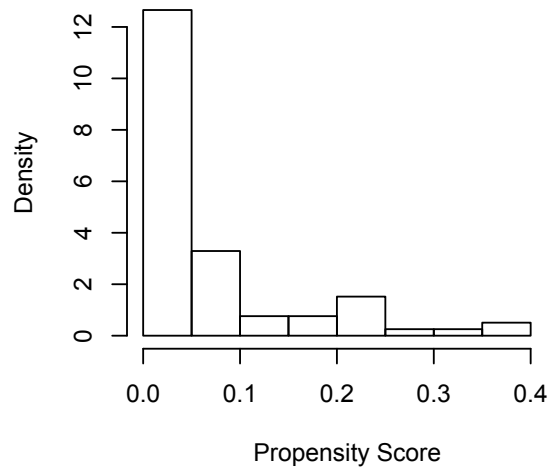
Matched Treated



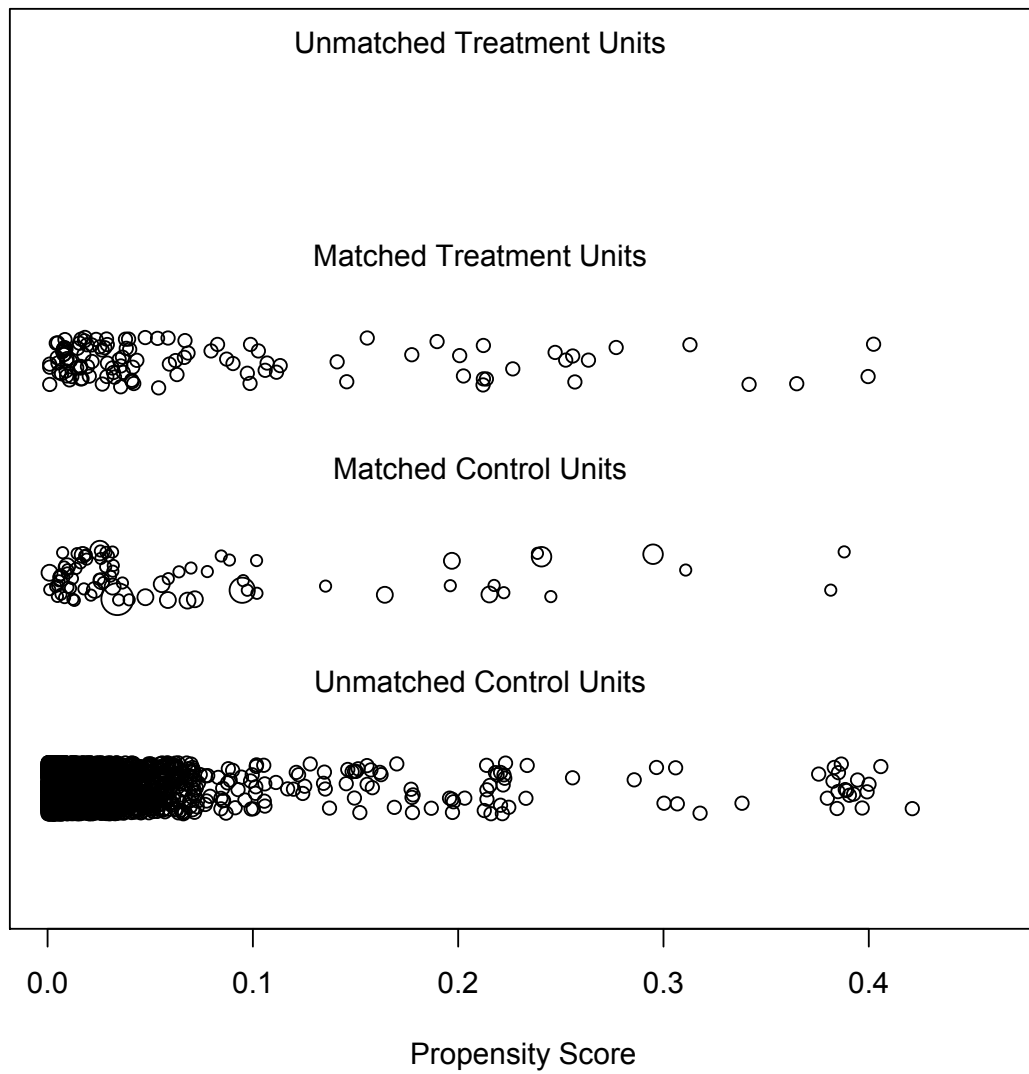
Raw Control



Matched Control



Distribution of Propensity Scores



FIRC: MATCHED ANALYSIS (with interwar, gdbuffer)

tab civwar10_2014 abdfirtarg, col chi2

civwar10_2014	abdfirtarg		Total
	0	1	
0	66 83.54	75 67.57	141 74.21
1	13 16.46	36 32.43	49 25.79
Total	79 100.00	111 100.00	190 100.00

Pearson chi2(1) = 6.1555 Pr = 0.013

tab civwar5_2014 abdfirtarg, col chi2

civwar5_2014	abdfirtarg		Total
	0	1	
0	71 89.87	81 72.97	152 80.00
1	8 10.13	30 27.03	38 20.00
Total	79 100.00	111 100.00	190 100.00

Pearson chi2(1) = 8.2390 Pr = 0.004

ttest civwar10_2014, by(abdfirtarg)

Two-sample t test with equal variances

Group	Obs	Mean	Std. Err.	Std. Dev.	[95% Conf. Interval]	
0	79	.164557	.0419826	.3731494	.080976	.2481379
1	111	.3243243	.0446337	.4702449	.2358709	.4127778
combined	190	.2578947	.0318217	.4386316	.1951234	.320666
diff		-.1597674	.0636799		-.2853863	-.0341485

diff = mean(0) - mean(1) t = -2.5089
Ho: diff = 0 degrees of freedom = 188

Ha: diff < 0 Pr(T < t) = 0.0065
Ha: diff != 0 Pr(|T| > |t|) = 0.0130
Ha: diff > 0 Pr(T > t) = 0.9935

ttest civwar5_2014, by(abdfirtarg)

Two-sample t test with equal variances

Group	Obs	Mean	Std. Err.	Std. Dev.	[95% Conf. Interval]	
0	79	.1012658	.0341586	.3036081	.0332613	.1692703
1	111	.2702703	.0423432	.4461134	.186356	.3541846
combined	190	.2	.0290957	.4010568	.1426059	.2573941
diff		-.1690044	.057894		-.2832097	-.0547992

diff = mean(0) - mean(1) t = -2.9192
Ho: diff = 0 degrees of freedom = 188

Ha: diff < 0 Pr(T < t) = 0.0020
Ha: diff != 0 Pr(|T| > |t|) = 0.0039
Ha: diff > 0 Pr(T > t) = 0.9980

LEADERSHIP FIRIC

Note: **peaceyrs** is omitted from matching because the balance between treated (32.11) and control (34.50) cases was already very good.

summary(leadfirc.out3)

Call:

```
matchit(formula = abdleaderfirc ~ lnpec4_fill + lntpop4_banks +
  mountainous + newstate + democracy + losewar10 + cw1000ongoing201411 +
  region1 + region2 + region3 + region4 + region5, data = leadfirc,
  method = "genetic", pop.size = 150)
```

Summary of balance for all data:

	Means Treated	Means Control	SD Control	Mean Diff	eQQ Med	eQQ Mean	eQQ Max
distance	0.0974	0.0069	0.0208	0.0905	0.0203	0.0878	0.3589
lnpec4_fill	4.3587	7.0217	3.8684	-2.6630	2.3751	2.6810	5.9940
lntpop4_banks	7.9191	8.5581	1.8705	-0.6390	0.7270	0.8145	3.1781
mountainous	0.2629	0.1832	0.2210	0.0797	0.0841	0.1035	0.3695
newstate	0.0469	0.0256	0.1580	0.0213	0.0000	0.0156	1.0000
democracy	0.1406	0.3557	0.4788	-0.2151	0.0000	0.2188	1.0000
losewar10	0.4688	0.0518	0.2217	0.4169	0.0000	0.4062	1.0000
cw1000ongoing201411	0.1562	0.0908	0.2873	0.0655	0.0000	0.0625	1.0000
region1	0.2969	0.2224	0.4159	0.0745	0.0000	0.0625	1.0000
region2	0.0156	0.1566	0.3635	-0.1410	0.0000	0.1406	1.0000
region3	0.1250	0.2046	0.4034	-0.0796	0.0000	0.0781	1.0000
region4	0.1719	0.1910	0.3931	-0.0192	0.0000	0.0312	1.0000
region5	0.3906	0.2253	0.4178	0.1653	0.0000	0.1562	1.0000

Summary of balance for matched data:

	Means Treated	Means Control	SD Control	Mean Diff	eQQ Med	eQQ Mean	eQQ Max
distance	0.0974	0.0858	0.1092	0.0117	0.0092	0.0401	0.2617
lnpec4_fill	4.3587	4.5584	3.8742	-0.1997	0.6931	1.2398	4.4886
lntpop4_banks	7.9191	8.2063	1.2563	-0.2873	0.6953	0.7835	2.1543
mountainous	0.2629	0.2411	0.1660	0.0217	0.0305	0.0406	0.3191
newstate	0.0469	0.0469	0.2137	0.0000	0.0000	0.0213	1.0000
democracy	0.1406	0.1406	0.3514	0.0000	0.0000	0.0213	1.0000
losewar10	0.4688	0.4688	0.5044	0.0000	0.0000	0.0851	1.0000
cw1000ongoing201411	0.1562	0.1875	0.3945	-0.0312	0.0000	0.0638	1.0000
region1	0.2969	0.2812	0.4545	0.0156	0.0000	0.0213	1.0000
region2	0.0156	0.0156	0.1254	0.0000	0.0000	0.0000	0.0000
region3	0.1250	0.1250	0.3343	0.0000	0.0000	0.0426	1.0000
region4	0.1719	0.1406	0.3514	0.0312	0.0000	0.0000	0.0000
region5	0.3906	0.4375	0.5014	-0.0469	0.0000	0.0638	1.0000

Percent Balance Improvement:

	Mean Diff.	eQQ Med	eQQ Mean	eQQ Max
distance	87.0943	54.5567	54.3738	27.0748
lnpec4_fill	92.5023	70.8164	53.7572	25.1140
lntpop4_banks	55.0445	4.3668	3.8009	32.2145
mountainous	72.7543	63.7564	60.8338	13.6309
newstate	100.0000	0.0000	-36.1702	0.0000
democracy	100.0000	0.0000	90.2736	0.0000
losewar10	100.0000	0.0000	79.0507	0.0000
cw1000ongoing201411	52.2777	0.0000	-2.1277	0.0000
region1	79.0172	0.0000	65.9574	0.0000
region2	100.0000	0.0000	100.0000	100.0000
region3	100.0000	0.0000	45.5319	0.0000
region4	-63.1300	0.0000	100.0000	100.0000
region5	71.6477	0.0000	59.1489	0.0000

Sample sizes:

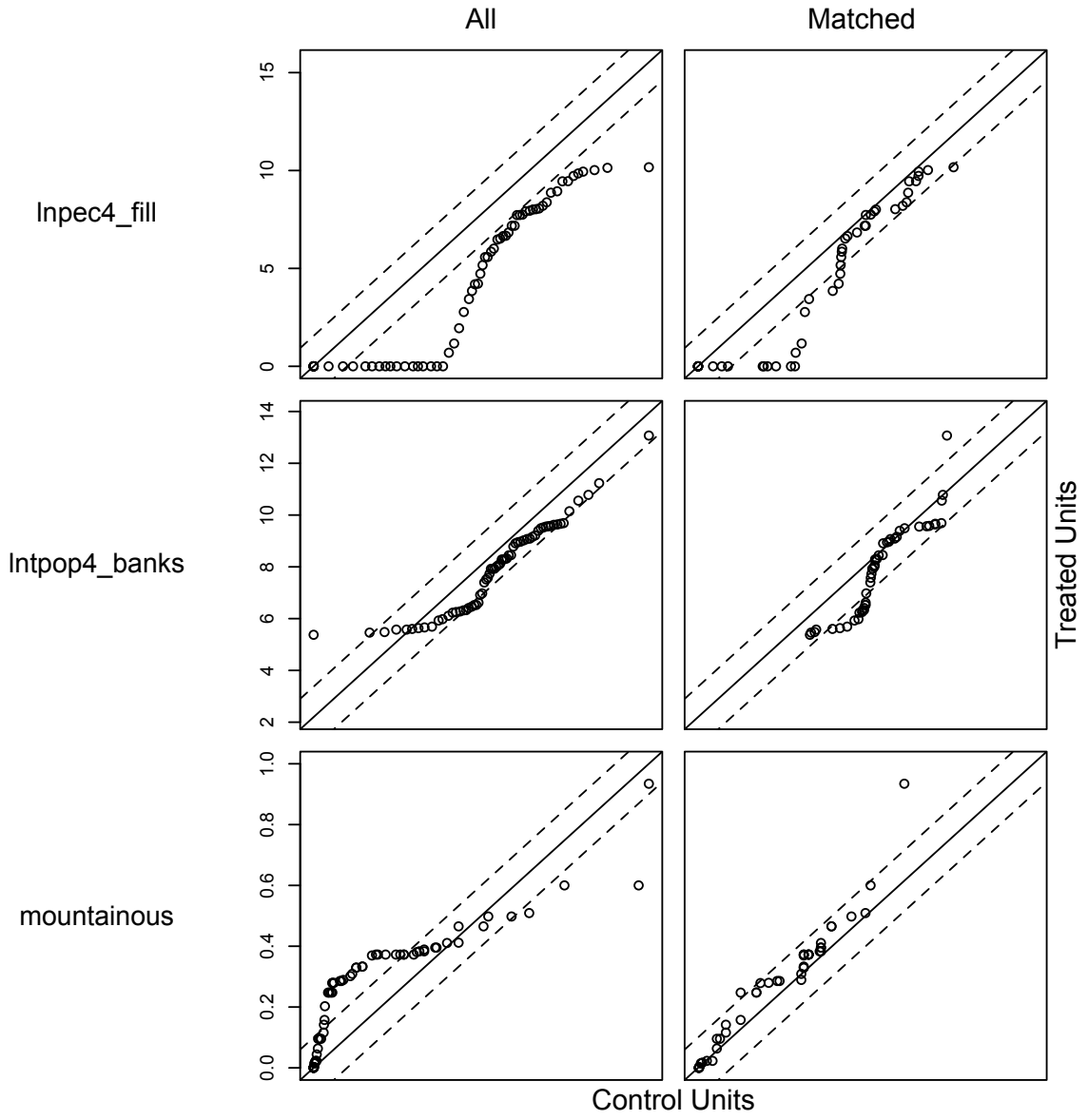
	Control	Treated
All	8318	64
Matched	47	64
Unmatched	8271	0
Discarded	0	0

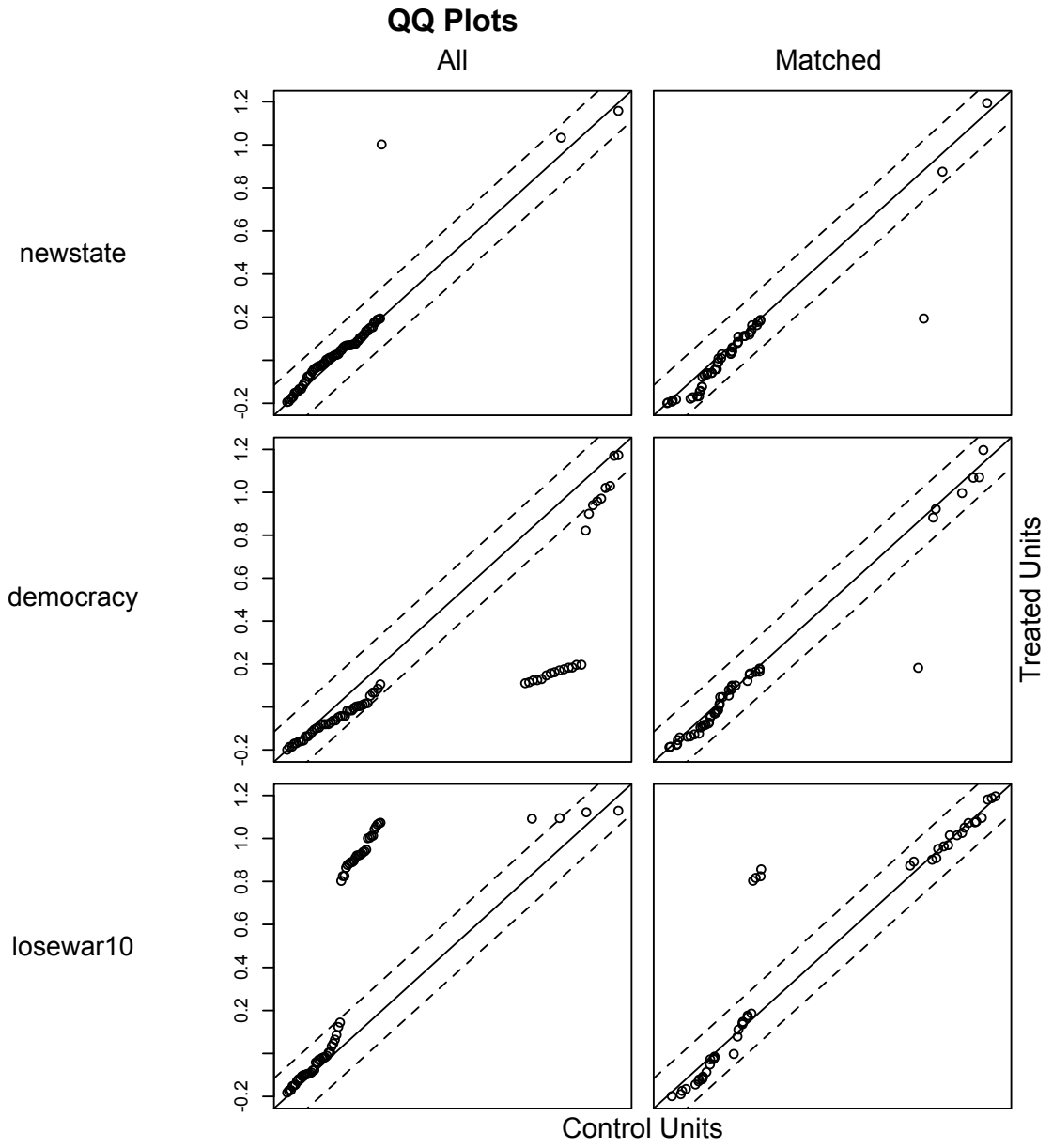
Table 4.5. Balance Statistics for Genetic Matching of Leadership Regime Change

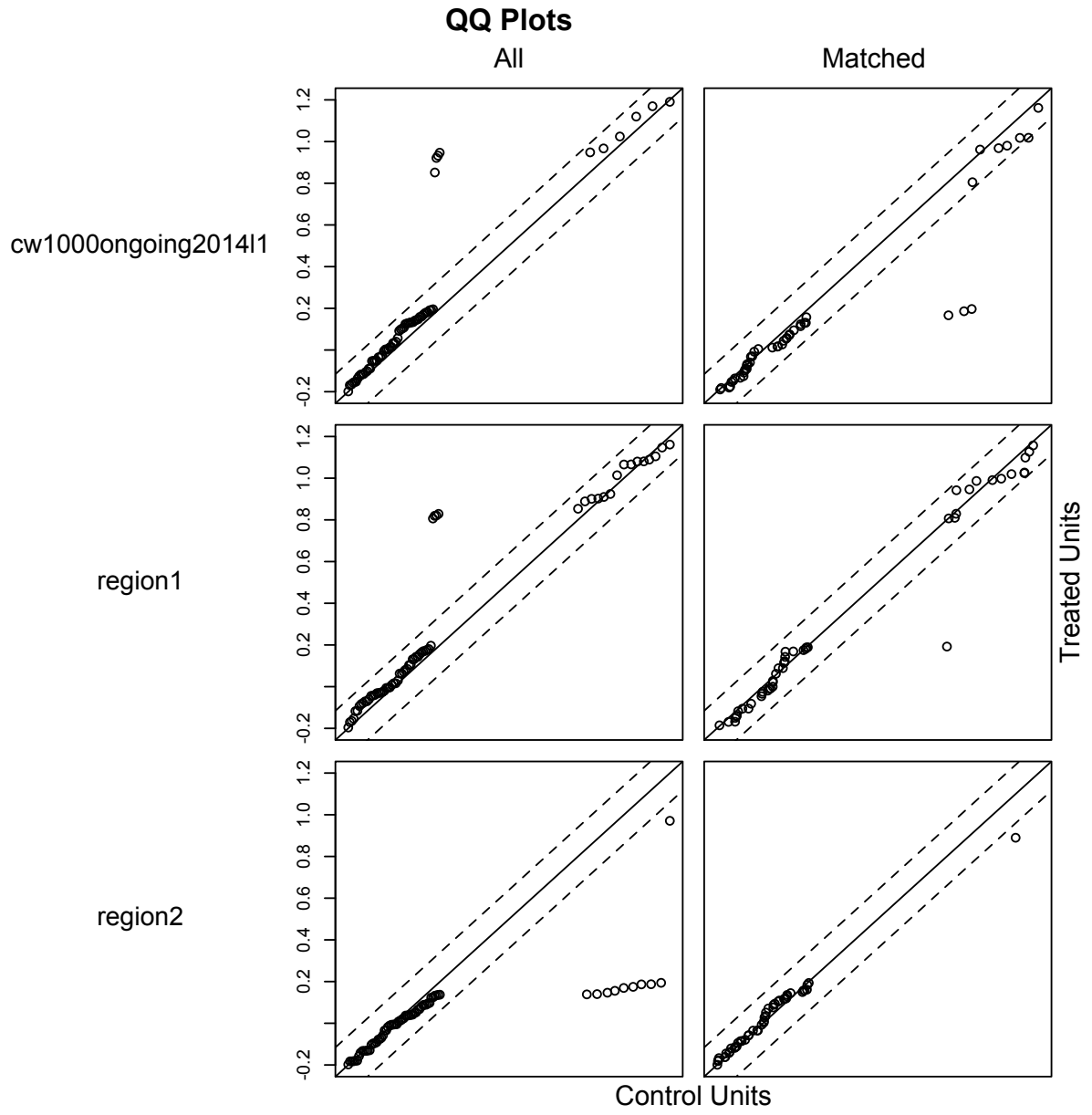
Variable	Mean Treated	Mean Control, before Matching	Mean Control, after Matching	Percent Improvement
Distance	0.0974	0.0069	0.0858	87.0943
Economic Development	4.3587	7.0217	4.5584	92.5023
Population	7.9191	8.5581	8.2063	55.0445
Rough Terrain	0.2629	0.1832	0.2411	72.7543
New State	0.0469	0.0256	0.0469	100
Democracy	0.1406	0.3557	0.1406	100
Lose Interstate War	0.4688	0.0518	0.4688	100
Ongoing Civil War	0.1562	0.0908	0.1875	52.2777
Peace Years	32.1094	34.4999	34.4999	NA
Europe	0.2969	0.2224	0.2812	79.0172
Middle East/North Africa	0.0156	0.1566	0.0156	100
Sub-Saharan Africa	0.1250	0.2046	0.1250	100
Asia	0.1719	0.1910	0.1406	-63.1300
Americas	0.3906	0.2253	0.4375	71.6477

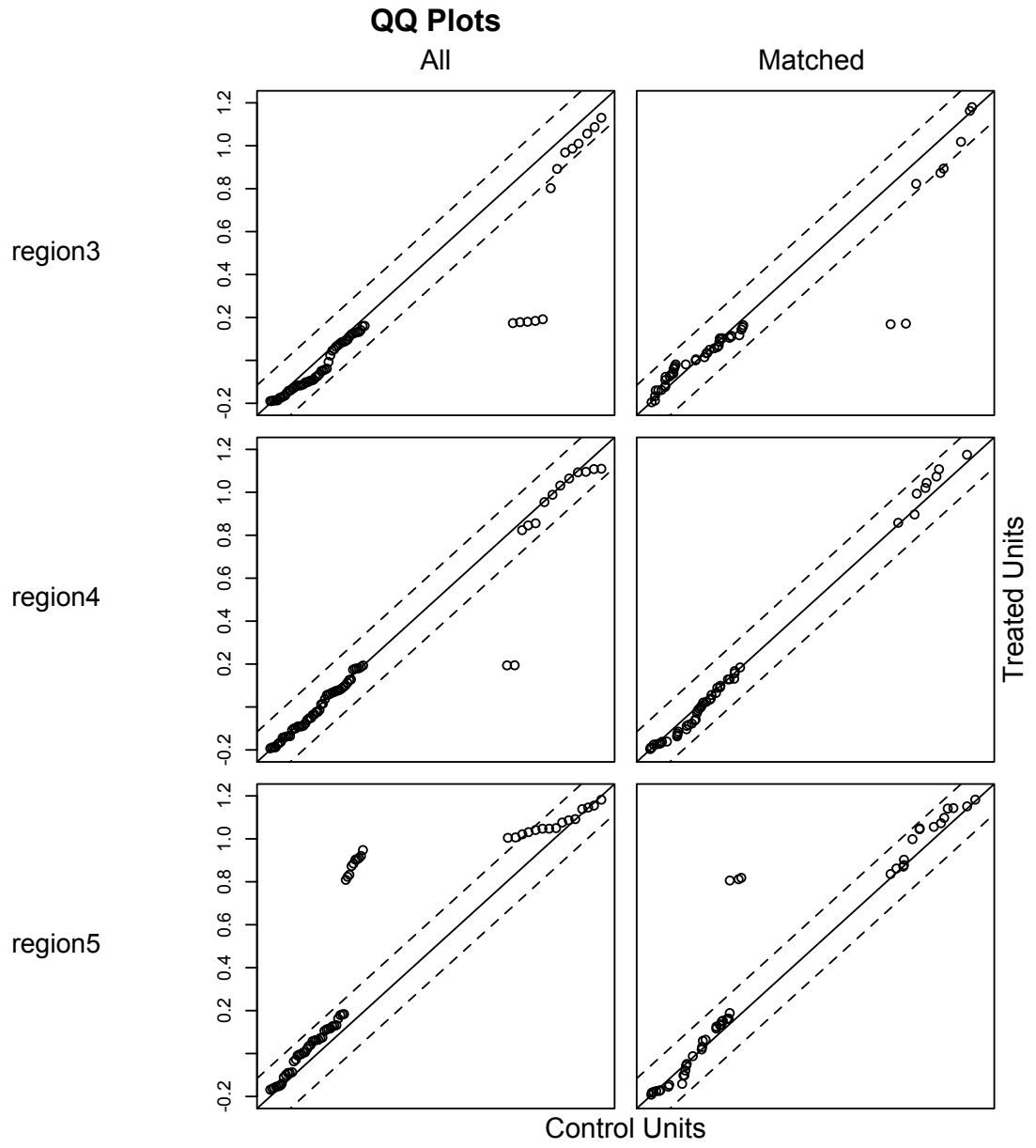
Note: *N* before matching was 8,382 (64 treated, 8,318 control). *N* after matching was 111 (64 treated, 47 control, some of which were matched to more than one case of regime change). The variable for peace years was omitted from matching because balance between treated and control cases was already acceptable and matching consistently worsened balance for this variable.

QQ Plots

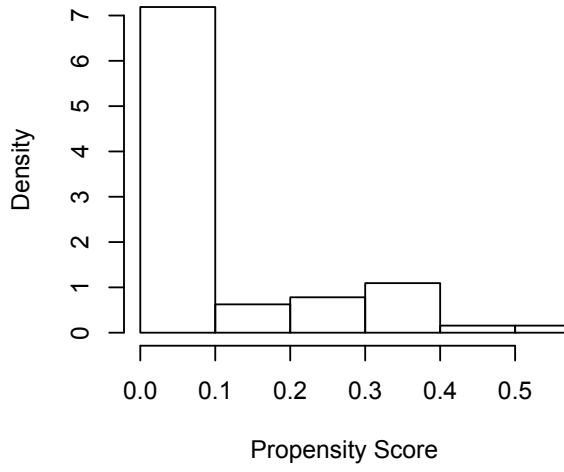




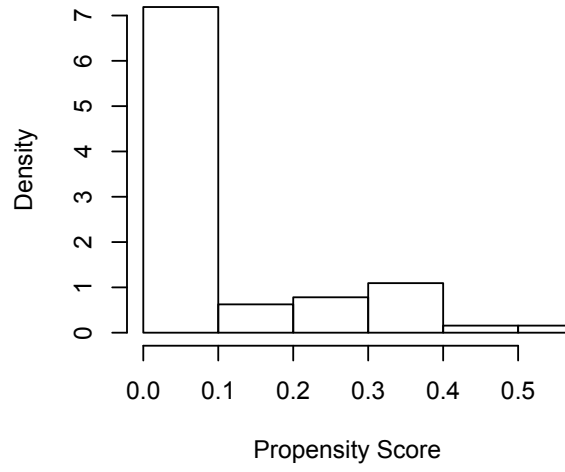




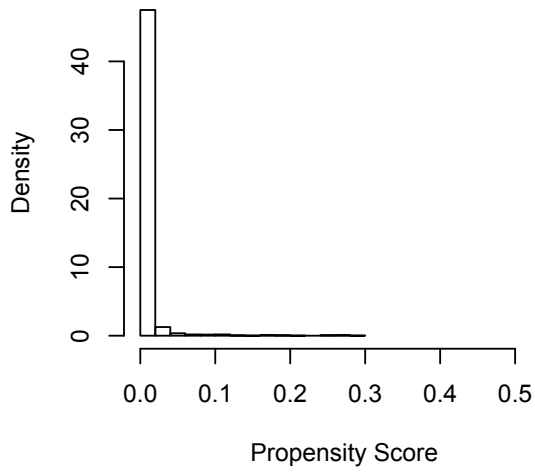
Raw Treated



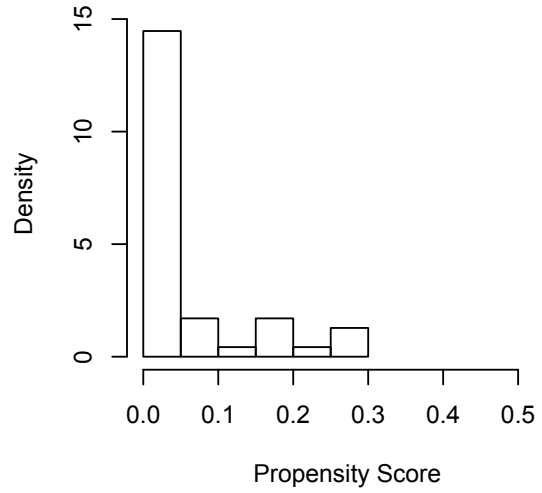
Matched Treated



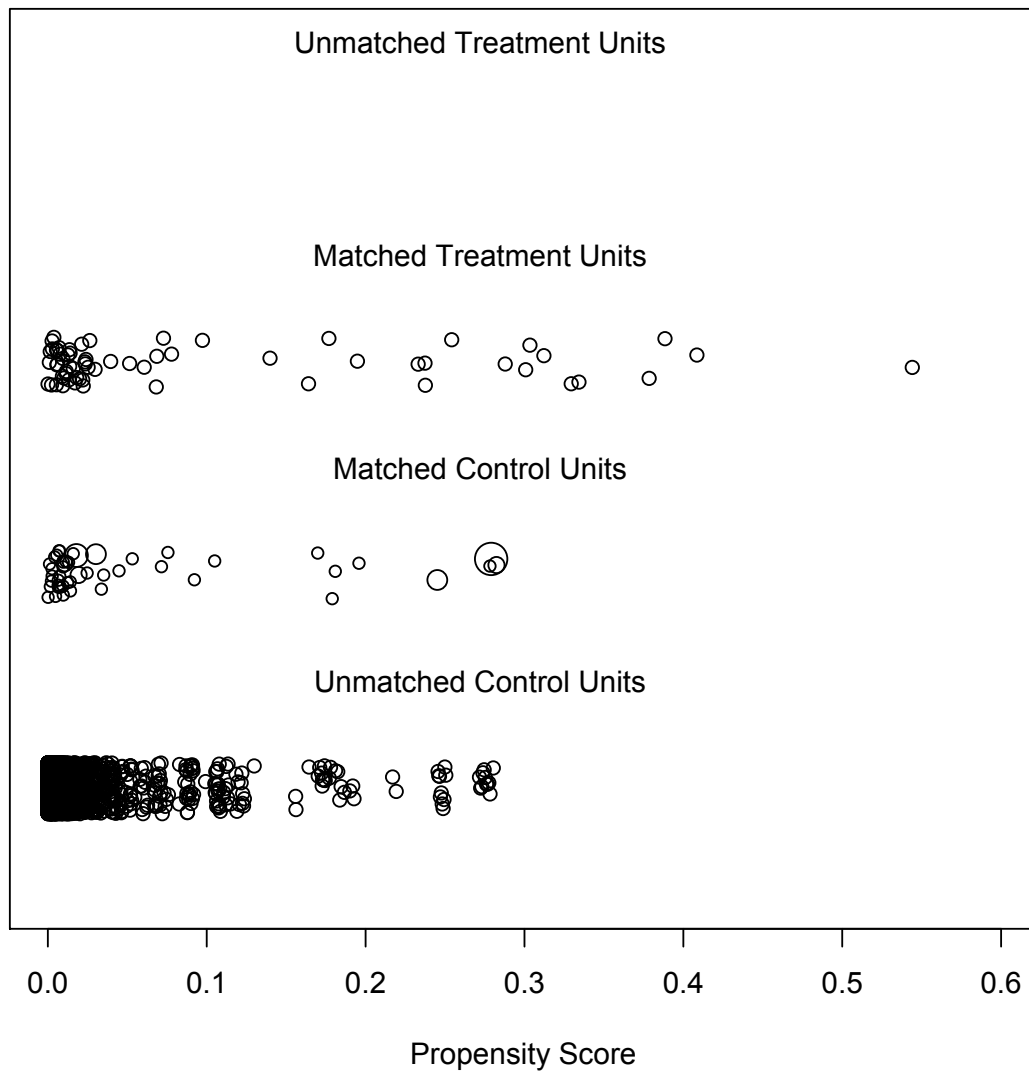
Raw Control



Matched Control



Distribution of Propensity Scores



LEADERSHIP FIRG: MATCHED ANALYSIS

tab civwar10_2014 abdleaderfirc, col chi2

civwar10_2014	abdleaderfirc		Total
	0	1	
0	37 78.72	34 53.12	71 63.96
1	10 21.28	30 46.88	40 36.04
Total	47 100.00	64 100.00	111 100.00

Pearson chi2(1) = 7.7039 Pr = 0.006

tab civwar5_2014 abdleaderfirc, col chi2

civwar5_2014	abdleaderfirc		Total
	0	1	
0	41 87.23	39 60.94	80 72.07
1	6 12.77	25 39.06	31 27.93
Total	47 100.00	64 100.00	111 100.00

Pearson chi2(1) = 9.3099 Pr = 0.002

ttest civwar10_2014, by(abdleaderfirc)

Two-sample t test with equal variances

Group	Obs	Mean	Std. Err.	Std. Dev.	[95% Conf. Interval]	
0	47	.212766	.0603426	.4136881	.0913026	.3342293
1	64	.46875	.0628709	.5029674	.3431125	.5943875
combined	111	.3603604	.0457762	.4822823	.2696427	.4510781
diff		-.255984	.0897816		-.4339283	-.0780397

diff = mean(0) - mean(1) t = -2.8512
Ho: diff = 0 degrees of freedom = 109

Ha: diff < 0 Ha: diff != 0 Ha: diff > 0
Pr(T < t) = 0.0026 Pr(|T| > |t|) = 0.0052 Pr(T > t) = 0.9974

ttest civwar5_2014, by(abdleaderfirc)

Two-sample t test with equal variances

Group	Obs	Mean	Std. Err.	Std. Dev.	[95% Conf. Interval]	
0	47	.1276596	.0492029	.3373181	.0286193	.2266999
1	64	.390625	.0614684	.4917474	.2677902	.5134598
combined	111	.2792793	.0427766	.4506797	.194506	.3640525
diff		-.2629654	.0832437		-.4279518	-.0979791

diff = mean(0) - mean(1) t = -3.1590
Ho: diff = 0 degrees of freedom = 109

Ha: diff < 0 Ha: diff != 0 Ha: diff > 0
Pr(T < t) = 0.0010 Pr(|T| > |t|) = 0.0020 Pr(T > t) = 0.9990

Rosenbaum Sensitivity Test for Leadership FIRC

```
#Load rbounds, which automatically loads Matching

library(rbounds)

Loading required package: Matching
Loading required package: MASS
##
## Matching (Version 4.9-5, Build Date: 2019-03-05)
## See http://sekhon.berkeley.edu/matching for additional documentation.
## Please cite software as:
## Jasjeet S. Sekhon. 2011. ``Multivariate and Propensity Score Matching
## Software with Automated Balance Optimization: The Matching package for R.''
## Journal of Statistical Software, 42(7): 1-52.
##

#Load rgenoud

library(rgenoud)
## rgenoud (Version 5.8-3.0, Build Date: 2019-01-22)
## See http://sekhon.berkeley.edu/rgenoud for additional documentation.
## Please cite software as:
## Walter Mebane, Jr. and Jasjeet S. Sekhon. 2011.
## ``Genetic Optimization Using Derivatives: The rgenoud package for R.''
## Journal of Statistical Software, 42(11): 1-26.

#Load foreign, which allows R to read in Stata files

library(foreign)

#Change directory to folder with data in it, then read in the data
#From "Miscellaneous" menu, choose "Change Working Directory," navigate to folder

lfirc=read.dta("leaderfirc_civwar_format12.dta")
attach(lfirc)
summary(lfirc)

#Specify the DV

Y <- lfirc$civwar10_2014

#Specify the Treatment

Tr <- lfirc$abdleaderfirc
#Specify the other variables

X <- cbind(lnpec4_fill, lntpop4_banks, mountainous, newstate, democracy, losewar10,
cwl000ongoing2014l1, gdbuffer, region1, region2, region3, region4, region5, peaceyrs)

#Specify the particular variables on which to match

BalanceMat <- cbind(lnpec4_fill, lntpop4_banks, mountainous, newstate, democracy, losewar10,
cwl000ongoing2014l1, gdbuffer, region1, region2, region3, region4, region5, peaceyrs)

#Genetic Weights (loads rgenoud automatically)

gen1 <- GenMatch(Tr=Tr, X=X, BalanceMat=BalanceMat, pop.size=50, data.type.int=FALSE,
replace=FALSE)

#Match

mgen1 <- Match(Y=Y, Tr=Tr, X=X, Weight.matrix=gen1, replace=FALSE)

#Show treatment effect

summary(mgen1)

Estimate... 0.26562
SE..... 0.067177
T-stat..... 3.9541
p.val..... 7.6827e-05
```

Original number of observations..... 8382
 Original number of treated obs..... 64
 Matched number of observations..... 64
 Matched number of observations (unweighted). 64

#Sensitivity Test

psens(mgen1, Gamma=4.0, GammaInc=.1)

Rosenbaum Sensitivity Test for Wilcoxon Signed Rank P-Value

Unconfounded estimate 2e-04

Gamma	Lower bound	Upper bound
1.0	2e-04	0.0002
1.1	1e-04	0.0004
1.2	0e+00	0.0009
1.3	0e+00	0.0016
1.4	0e+00	0.0027
1.5	0e+00	0.0042
1.6	0e+00	0.0061
1.7	0e+00	0.0086
1.8	0e+00	0.0116
1.9	0e+00	0.0153
2.0	0e+00	0.0195
2.1	0e+00	0.0243
2.2	0e+00	0.0298
2.3	0e+00	0.0358
2.4	0e+00	0.0425
2.5	0e+00	0.0496
2.6	0e+00	0.0573
2.7	0e+00	0.0655
2.8	0e+00	0.0742
2.9	0e+00	0.0832
3.0	0e+00	0.0927
3.1	0e+00	0.1025
3.2	0e+00	0.1127
3.3	0e+00	0.1232
3.4	0e+00	0.1339
3.5	0e+00	0.1448
3.6	0e+00	0.1560
3.7	0e+00	0.1673
3.8	0e+00	0.1788
3.9	0e+00	0.1904
4.0	0e+00	0.2021

Note: Gamma is Odds of Differential Assignment To Treatment Due to Unobserved Factors

LEADERSHIP FIRIC (with interwar and gdbuffer)

summary(leadfirc.out4)

Call:

```
matchit(formula = abdleaderfirc ~ lnpec4_fill + lntpop4_banks +
  mountainous + newstate + democracy + interwar + cw1000ongoing201411 +
  gdbuffer + region1 + region2 + region3 + region4 + region5,
  data = leadfirc, method = "genetic", pop.size = 150)
```

Summary of balance for all data:

	Means	Treated	Means	Control	SD	Control	Mean Diff	eQQ	Med	eQQ	Mean	eQQ	Max
distance	0.0766		0.0071	0.0217	0.0695	0.0258	0.0666	0.2726					
lnpec4_fill	4.3587		7.0217	3.8684	-2.6630	2.3751	2.6810	5.9940					
lntpop4_banks	7.9191		8.5581	1.8705	-0.6390	0.7270	0.8145	3.1781					
mountainous	0.2629		0.1832	0.2210	0.0797	0.0841	0.1035	0.3695					
newstate	0.0469		0.0256	0.1580	0.0213	0.0000	0.0156	1.0000					
democracy	0.1406		0.3557	0.4788	-0.2151	0.0000	0.2188	1.0000					
interwar	0.4531		0.0513	0.2207	0.4018	0.0000	0.3906	1.0000					
cw1000ongoing201411	0.1562		0.0908	0.2873	0.0655	0.0000	0.0625	1.0000					
gdbuffer	0.2812		0.1193	0.3241	0.1620	0.0000	0.1562	1.0000					
region1	0.2969		0.2224	0.4159	0.0745	0.0000	0.0625	1.0000					
region2	0.0156		0.1566	0.3635	-0.1410	0.0000	0.1406	1.0000					
region3	0.1250		0.2046	0.4034	-0.0796	0.0000	0.0781	1.0000					
region4	0.1719		0.1910	0.3931	-0.0192	0.0000	0.0312	1.0000					
region5	0.3906		0.2253	0.4178	0.1653	0.0000	0.1562	1.0000					

Summary of balance for matched data:

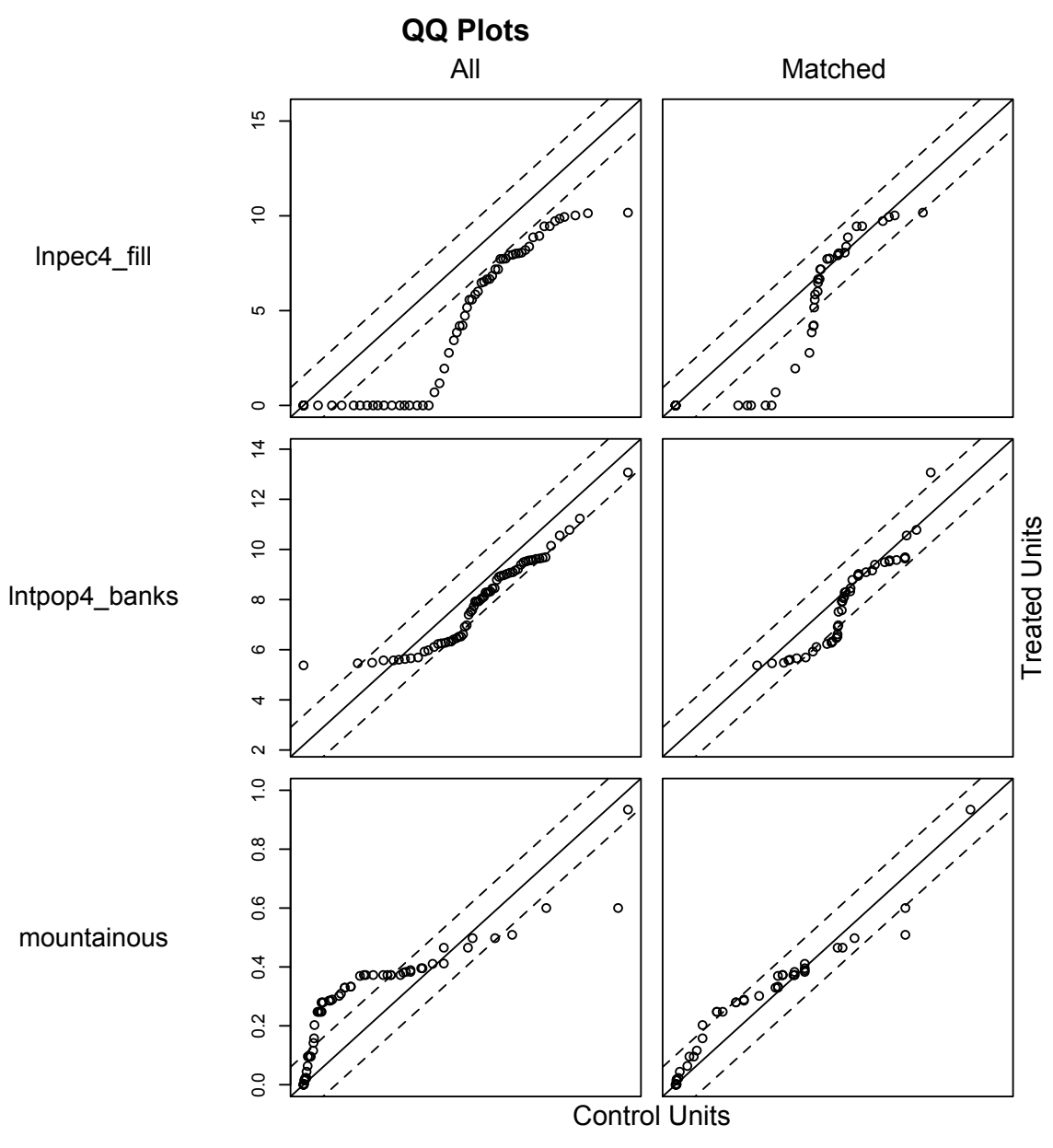
	Means	Treated	Means	Control	SD	Control	Mean Diff	eQQ	Med	eQQ	Mean	eQQ	Max
distance	0.0766		0.0739	0.0962	0.0026	0.0124	0.0163	0.0673					
lnpec4_fill	4.3587		4.4592	3.8517	-0.1005	0.3060	1.0752	4.5951					
lntpop4_banks	7.9191		8.0897	1.5001	-0.1707	0.4942	0.6457	1.6389					
mountainous	0.2629		0.2491	0.2119	0.0137	0.0293	0.0427	0.1991					
newstate	0.0469		0.0469	0.2137	0.0000	0.0000	0.0217	1.0000					
democracy	0.1406		0.1406	0.3515	0.0000	0.0000	0.0000	0.0000					
interwar	0.4531		0.4531	0.5033	0.0000	0.0000	0.0652	1.0000					
cw1000ongoing201411	0.1562		0.1562	0.3671	0.0000	0.0000	0.0217	1.0000					
gdbuffer	0.2812		0.2656	0.4465	0.0156	0.0000	0.0217	1.0000					
region1	0.2969		0.2969	0.4619	0.0000	0.0000	0.0435	1.0000					
region2	0.0156		0.0156	0.1254	0.0000	0.0000	0.0000	0.0000					
region3	0.1250		0.1250	0.3344	0.0000	0.0000	0.0217	1.0000					
region4	0.1719		0.1562	0.3671	0.0156	0.0000	0.0435	1.0000					
region5	0.3906		0.4062	0.4966	-0.0156	0.0000	0.1087	1.0000					

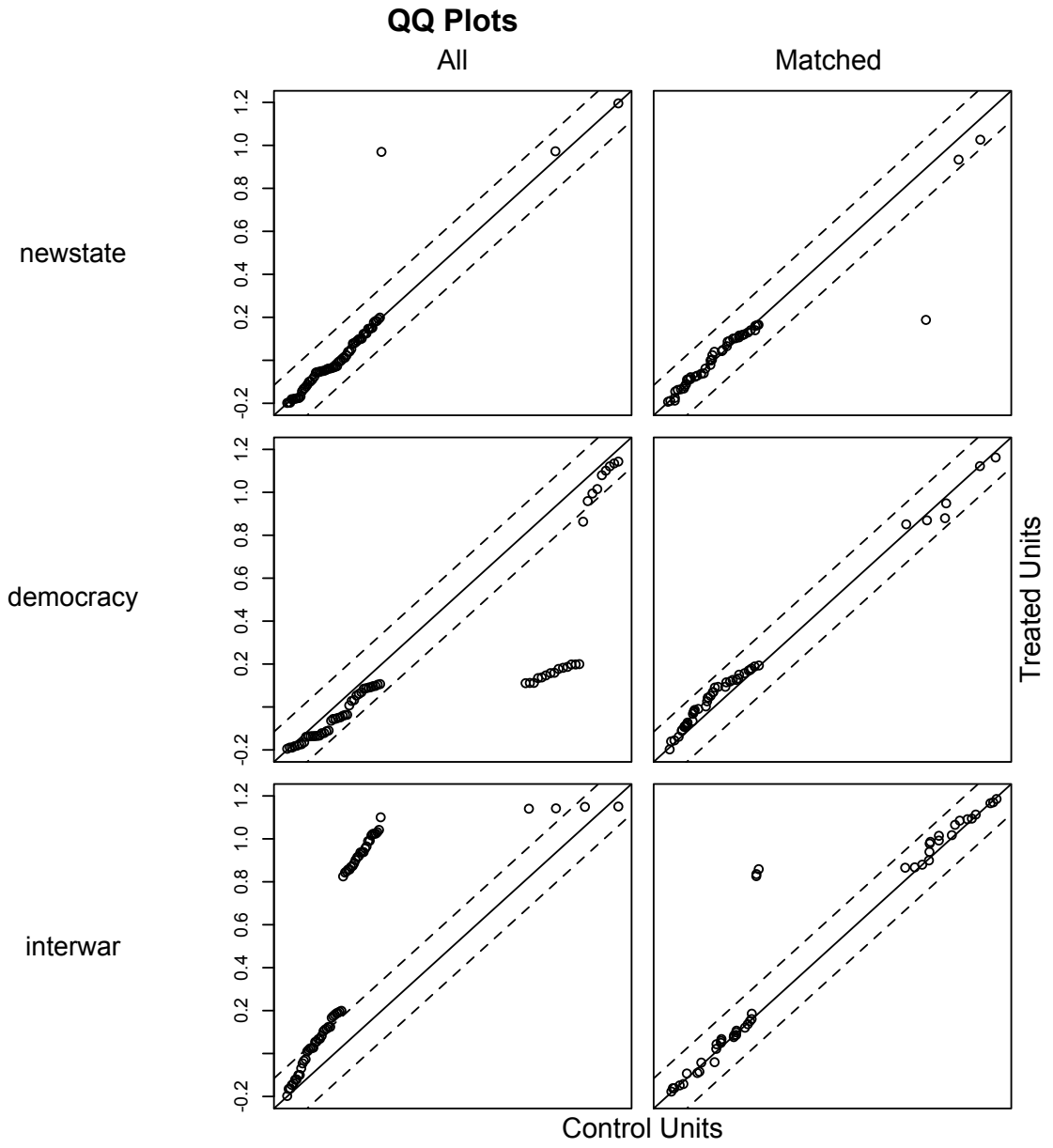
Percent Balance Improvement:

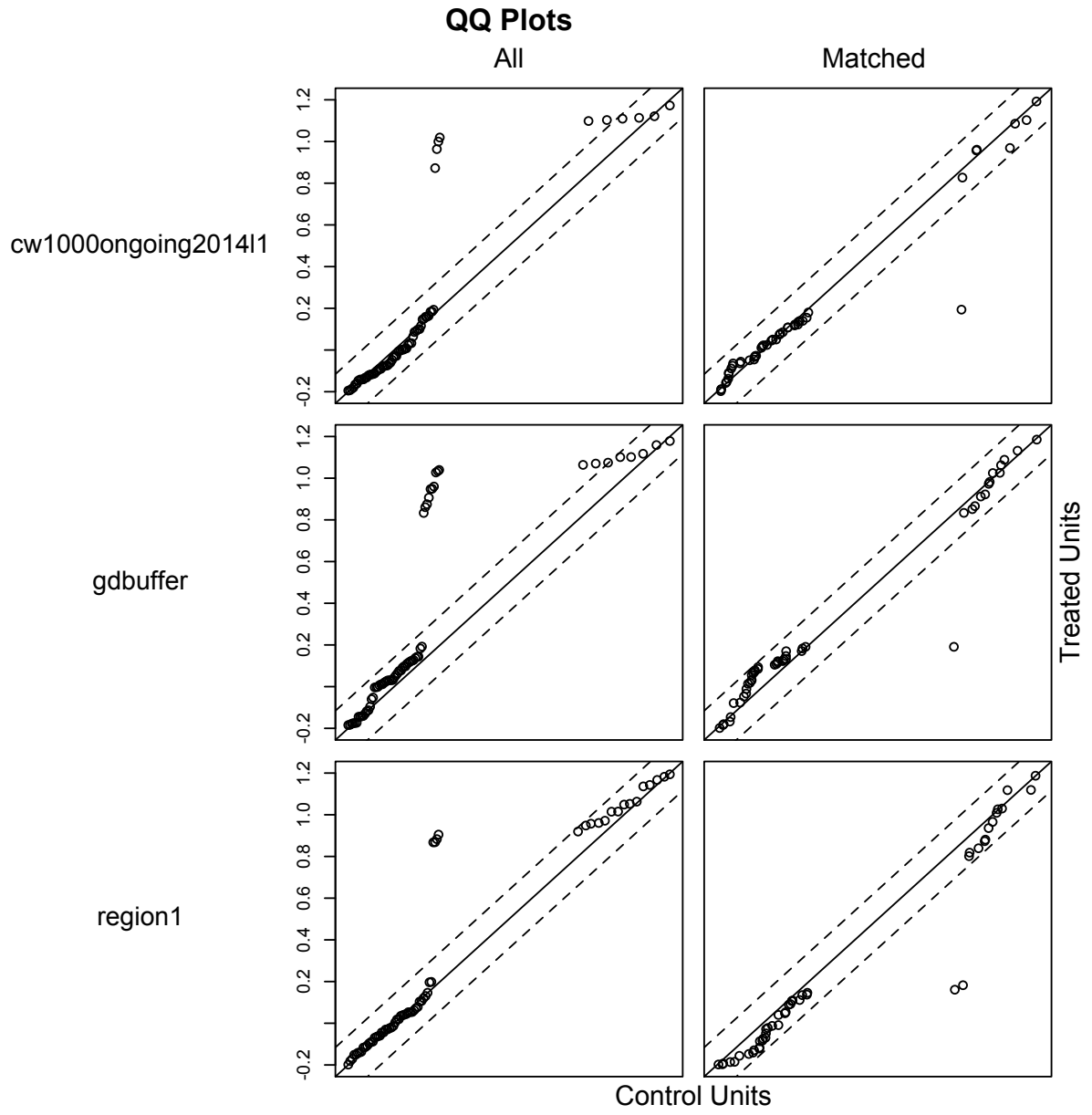
	Mean Diff.	eQQ	Med	eQQ	Mean	eQQ	Max
distance	96.1925	51.9342	75.5507	75.3198			
lnpec4_fill	96.2263	87.1156	59.8970	23.3375			
lntpop4_banks	73.2930	32.0264	20.7182	48.4307			
mountainous	82.7524	65.1800	58.7313	46.1189			
newstate	100.0000	0.0000	-39.1304	0.0000			
democracy	100.0000	0.0000	100.0000	100.0000			
interwar	100.0000	0.0000	83.3043	0.0000			
cw1000ongoing201411	100.0000	0.0000	65.2174	0.0000			
gdbuffer	90.3544	0.0000	86.0870	0.0000			
region1	100.0000	0.0000	30.4348	0.0000			
region2	100.0000	0.0000	100.0000	100.0000			
region3	100.0000	0.0000	72.1739	0.0000			
region4	18.4350	0.0000	-39.1304	0.0000			
region5	90.5492	0.0000	30.4348	0.0000			

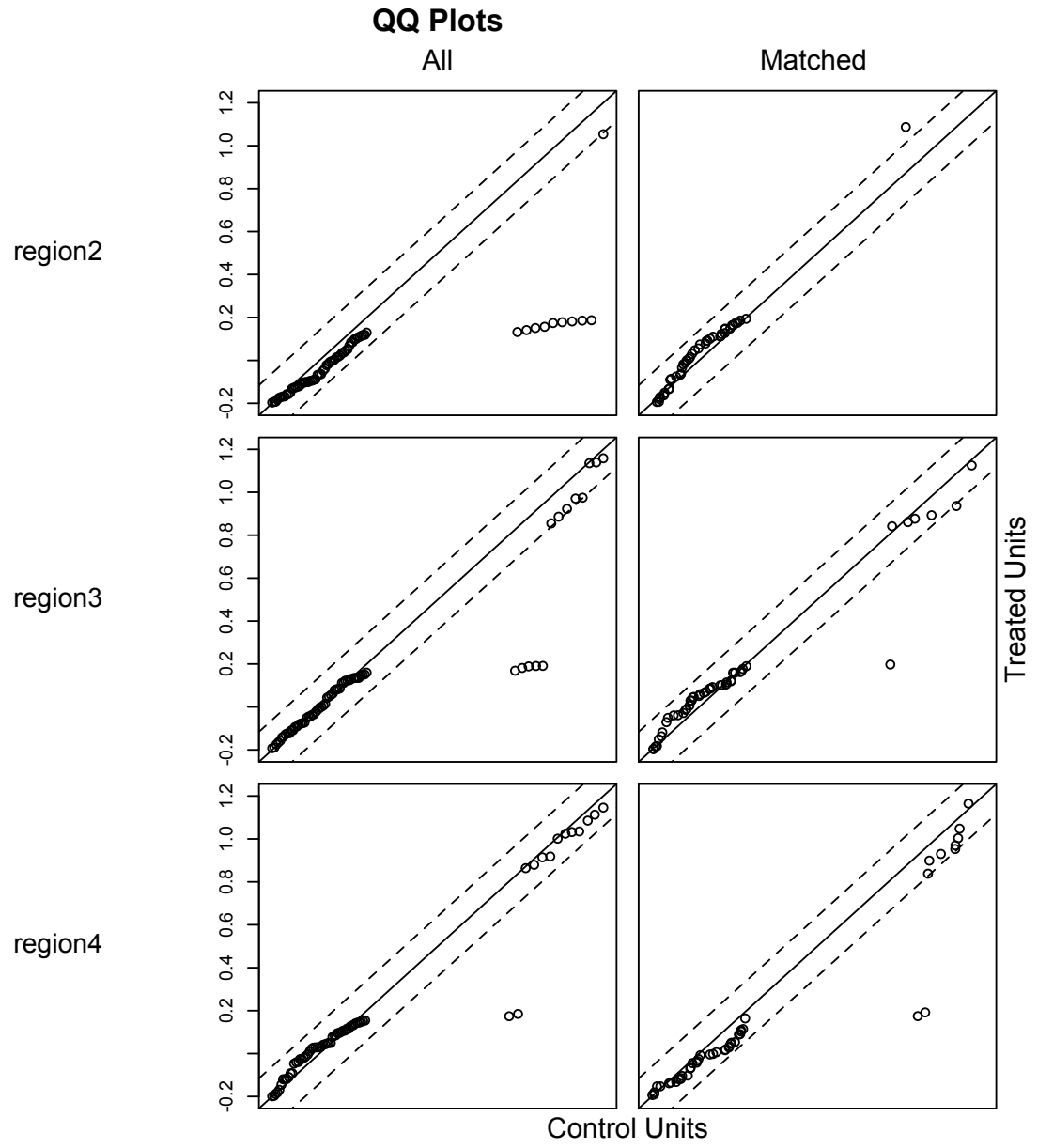
Sample sizes:

	Control	Treated
All	8318	64
Matched	46	64
Unmatched	8272	0
Discarded	0	0

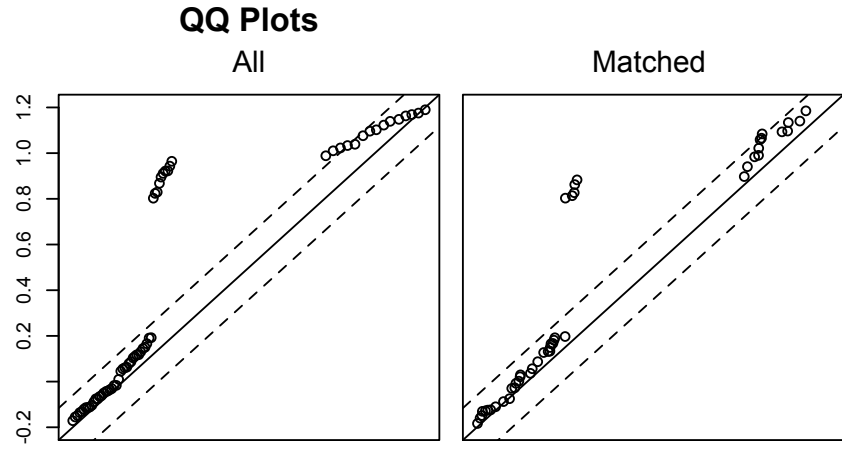








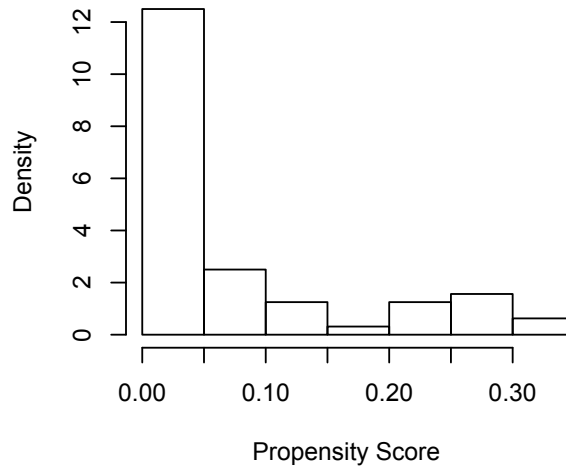
region5



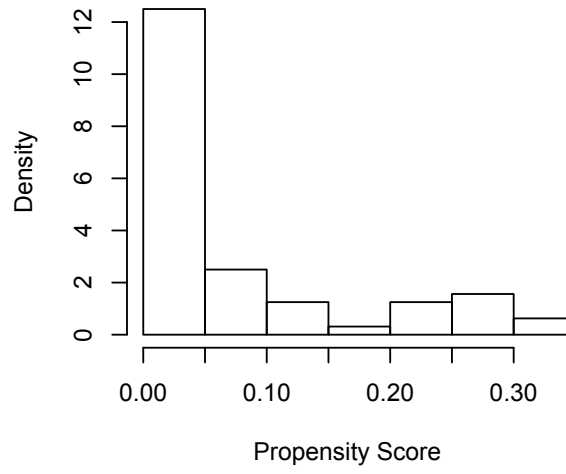
Treated Units

Control Units

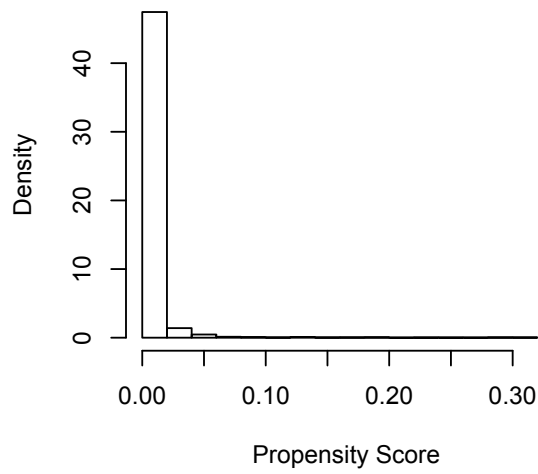
Raw Treated



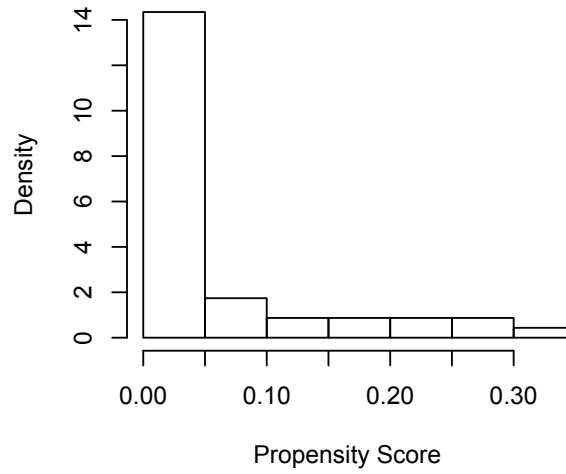
Matched Treated



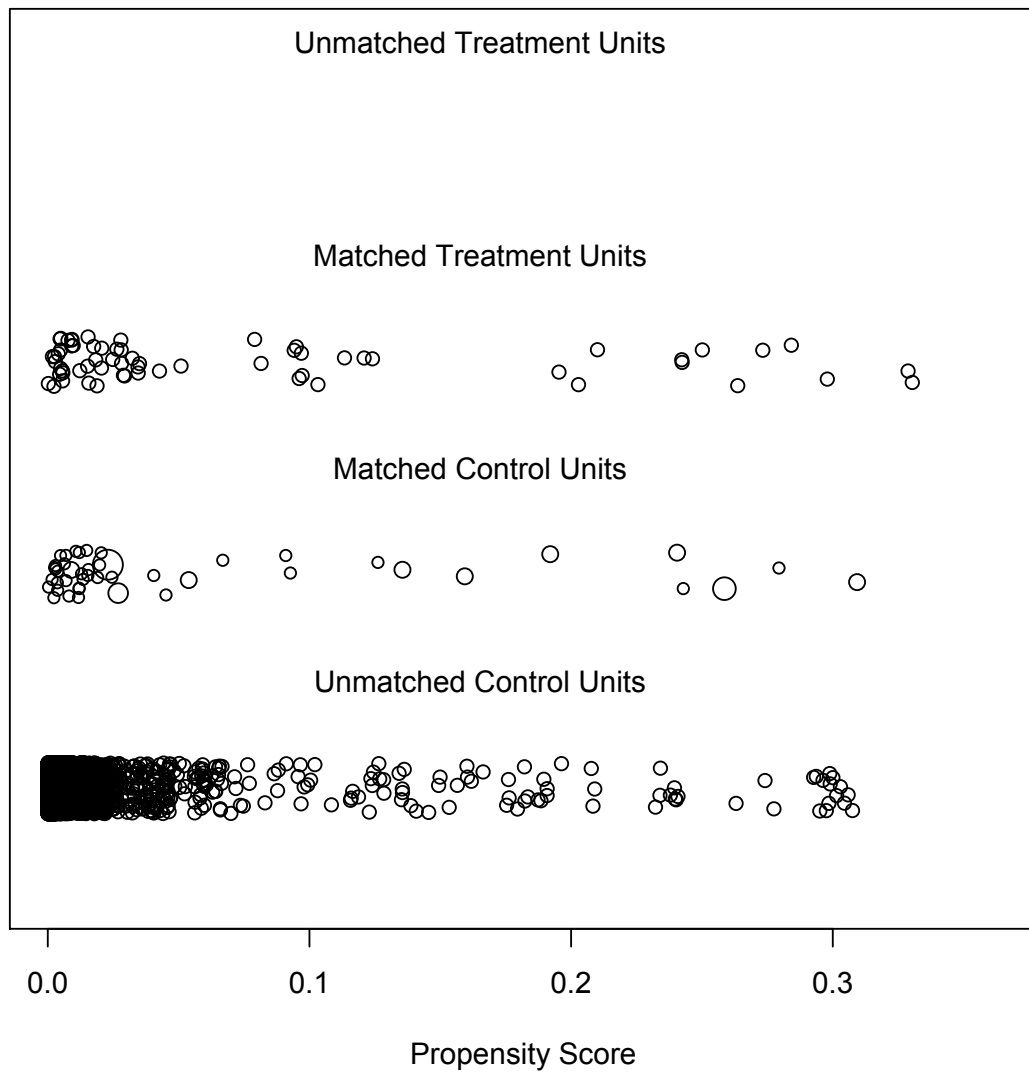
Raw Control



Matched Control



Distribution of Propensity Scores



LEADERSHIP FIRG: MATCHED ANALYSIS (with interwar and gdbuffer)

tab civwar10_2014 abdleaderfirc, col chi2

civwar10_2014	abdleaderfirc		Total
	0	1	
0	36 78.26	34 53.12	70 63.64
1	10 21.74	30 46.88	40 36.36
Total	46 100.00	64 100.00	110 100.00

Pearson chi2(1) = 7.3074 Pr = 0.007

tab civwar5_2014 abdleaderfirc, col chi2

civwar5_2014	abdleaderfirc		Total
	0	1	
0	40 86.96	39 60.94	79 71.82
1	6 13.04	25 39.06	31 28.18
Total	46 100.00	64 100.00	110 100.00

Pearson chi2(1) = 8.9521 Pr = 0.003

ttest civwar10_2014, by(abdleaderfirc)

Two-sample t test with equal variances

Group	Obs	Mean	Std. Err.	Std. Dev.	[95% Conf. Interval]	
0	46	.2173913	.0614875	.4170288	.093549	.3412336
1	64	.46875	.0628709	.5029674	.3431125	.5943875
combined	110	.3636364	.0460758	.4832473	.2723156	.4549571
diff		-.2513587	.0906717		-.4310857	-.0716317

diff = mean(0) - mean(1) t = -2.7722
Ho: diff = 0 degrees of freedom = 108

Ha: diff < 0 Pr(T < t) = 0.0033
Ha: diff != 0 Pr(|T| > |t|) = 0.0066
Ha: diff > 0 Pr(T > t) = 0.9967

ttest civwar5_2014, by(abdleaderfirc)

Two-sample t test with equal variances

Group	Obs	Mean	Std. Err.	Std. Dev.	[95% Conf. Interval]	
0	46	.1304348	.0502044	.3405026	.029318	.2315516
1	64	.390625	.0614684	.4917474	.2677902	.5134598
combined	110	.2818182	.0430912	.4519442	.1964128	.3672235
diff		-.2601902	.0841165		-.4269236	-.0934568

diff = mean(0) - mean(1) t = -3.0932
Ho: diff = 0 degrees of freedom = 108

Ha: diff < 0 Pr(T < t) = 0.0013
Ha: diff != 0 Pr(|T| > |t|) = 0.0025
Ha: diff > 0 Pr(T > t) = 0.9987

INSTITUTIONAL FIRIC

summary(instfirc.out2)

Note: **peaceyrs** is omitted from matching because the balance between treated (34.15) and control (34.48) cases was already very good.

Call:

```
matchit(formula = abdstfirc ~ lnpec4_fill + lntpop4_banks +
  mountainous + newstate + democracy + losewar10 + cw1000ongoing201411 +
  region1 + region2 + region3 + region4 + region5, data = instfirc,
  method = "genetic", pop.size = 150)
```

Summary of balance for all data:

	Means Treated	Means Control	SD Control	Mean Diff	eQQ Med	eQQ Mean	eQQ Max
distance	0.0164	0.0024	0.0067	0.0140	0.0073	0.0200	0.1531
lnpec4_fill	5.8619	7.0217	3.8684	-1.1598	0.7282	1.2457	4.4543
lntpop4_banks	8.1346	8.5581	1.8705	-0.4234	0.5903	0.7358	2.7548
mountainous	0.1615	0.1832	0.2210	-0.0216	0.0371	0.0704	0.5023
newstate	0.1000	0.0256	0.1580	0.0744	0.0000	0.0500	1.0000
democracy	0.0500	0.3557	0.4788	-0.3057	0.0000	0.3000	1.0000
losewar10	0.3000	0.0518	0.2217	0.2482	0.0000	0.2500	1.0000
cw1000ongoing201411	0.0500	0.0908	0.2873	-0.0408	0.0000	0.0500	1.0000
region1	0.4500	0.2224	0.4159	0.2276	0.0000	0.2000	1.0000
region2	0.0500	0.1566	0.3635	-0.1066	0.0000	0.1000	1.0000
region3	0.0000	0.2046	0.4034	-0.2046	0.0000	0.2000	1.0000
region4	0.1000	0.1910	0.3931	-0.0910	0.0000	0.1000	1.0000
region5	0.4000	0.2253	0.4178	0.1747	0.0000	0.1500	1.0000

Summary of balance for matched data:

	Means Treated	Means Control	SD Control	Mean Diff	eQQ Med	eQQ Mean	eQQ Max
distance	0.0164	0.0164	0.0188	0.0000	0.0003	0.0027	0.0245
lnpec4_fill	5.8619	5.7760	4.5245	0.0859	0.5857	0.7685	3.5484
lntpop4_banks	8.1346	8.1731	1.9432	-0.0384	0.3405	0.3916	1.0523
mountainous	0.1615	0.1621	0.1528	-0.0006	0.0178	0.0304	0.0883
newstate	0.1000	0.1000	0.3082	0.0000	0.0000	0.0526	1.0000
democracy	0.0500	0.0500	0.2239	0.0000	0.0000	0.0000	0.0000
losewar10	0.3000	0.3000	0.4708	0.0000	0.0000	0.0526	1.0000
cw1000ongoing201411	0.0500	0.0500	0.2239	0.0000	0.0000	0.0000	0.0000
region1	0.4500	0.4500	0.5111	0.0000	0.0000	0.0526	1.0000
region2	0.0500	0.0500	0.2239	0.0000	0.0000	0.0000	0.0000
region3	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
region4	0.1000	0.1000	0.3082	0.0000	0.0000	0.0526	1.0000
region5	0.4000	0.4000	0.5033	0.0000	0.0000	0.0000	0.0000

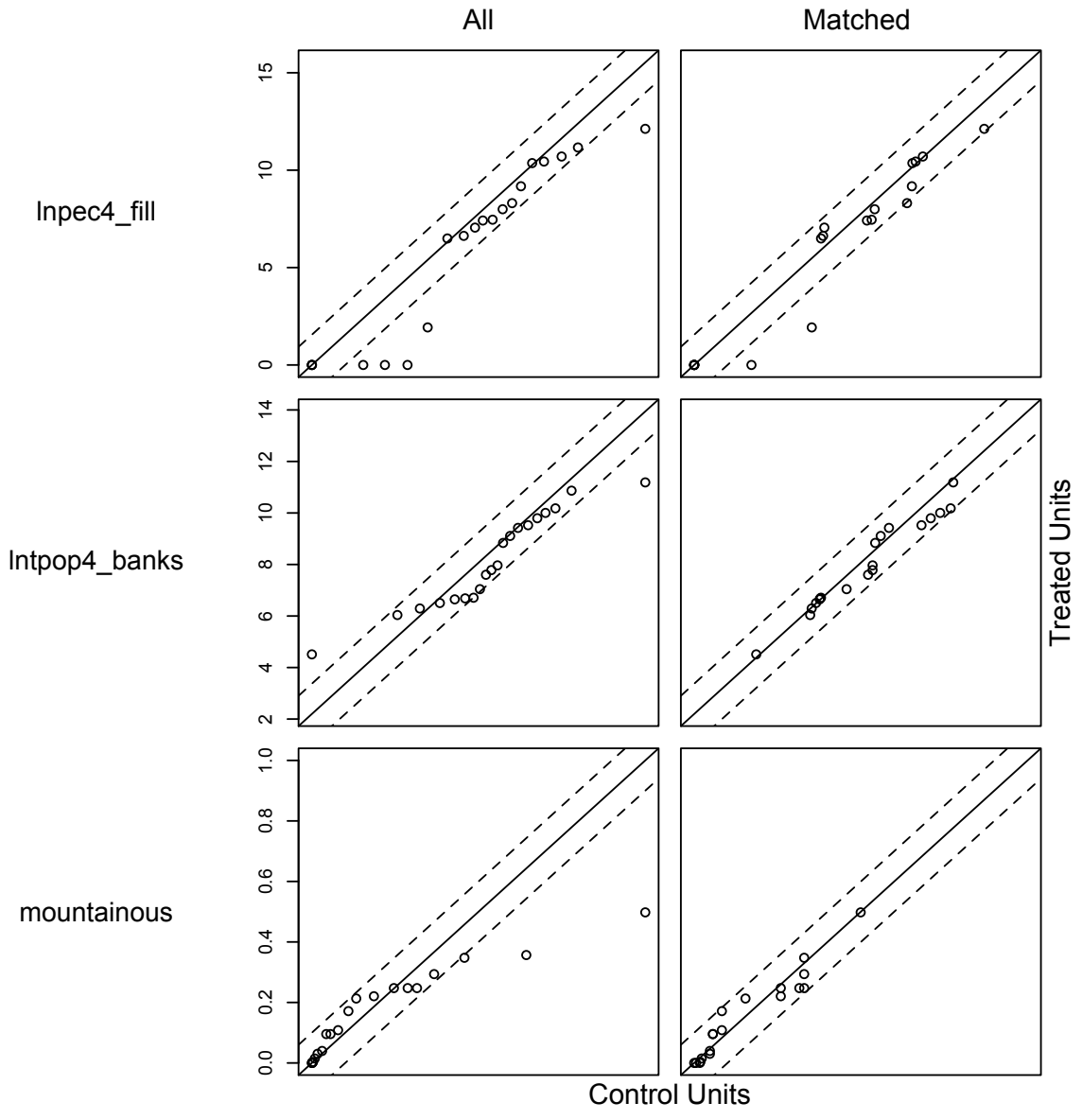
Percent Balance Improvement:

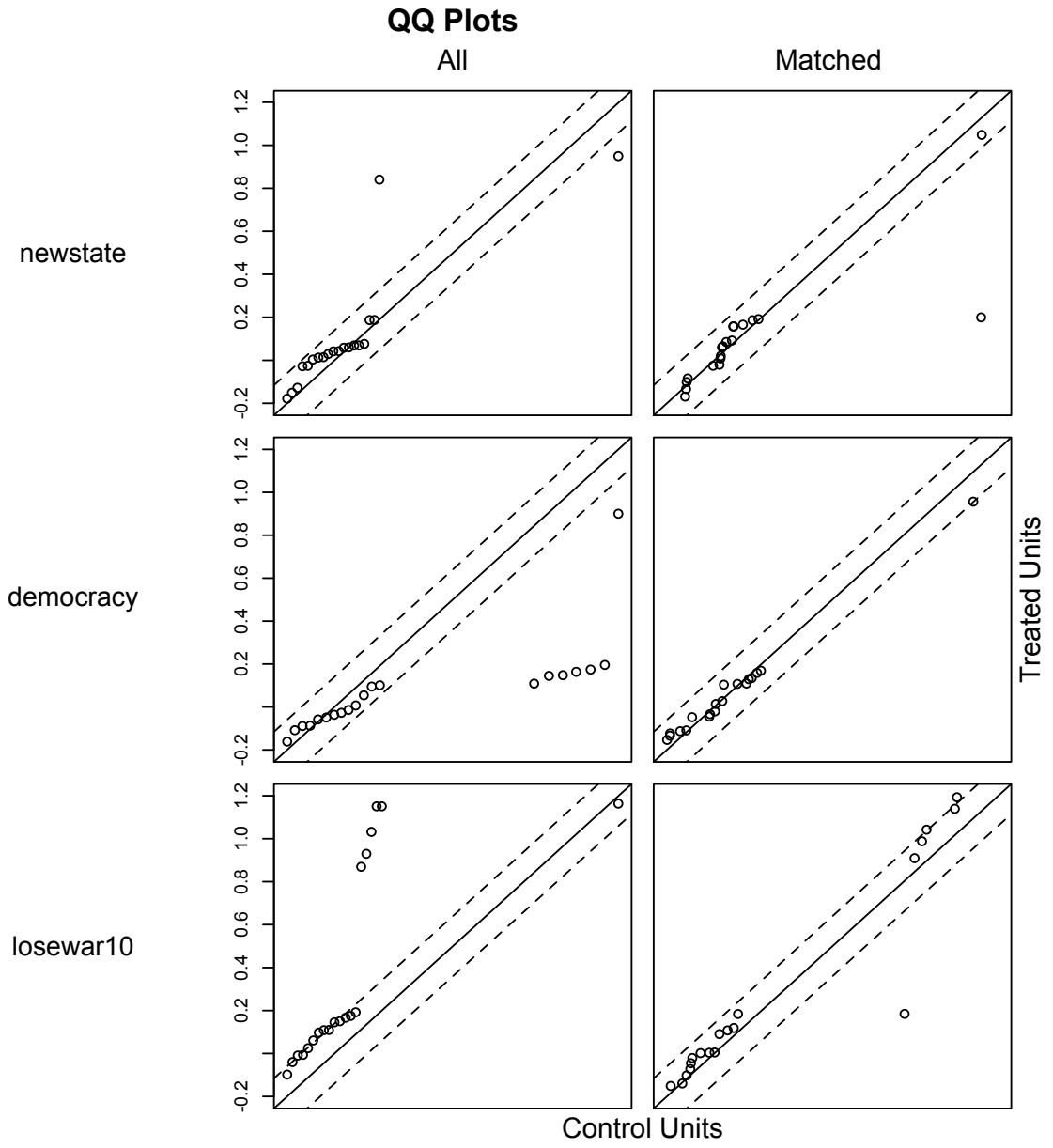
	Mean Diff.	eQQ Med	eQQ Mean	eQQ Max
distance	99.6527	95.3279	86.4497	84.0051
lnpec4_fill	92.5948	19.5730	38.3051	20.3383
lntpop4_banks	90.9256	42.3236	46.7759	61.8011
mountainous	97.3215	51.9136	56.8106	82.4201
newstate	100.0000	0.0000	-5.2632	0.0000
democracy	100.0000	0.0000	100.0000	100.0000
losewar10	100.0000	0.0000	78.9474	0.0000
cw1000ongoing201411	100.0000	0.0000	100.0000	100.0000
region1	100.0000	0.0000	73.6842	0.0000
region2	100.0000	0.0000	100.0000	100.0000
region3	100.0000	0.0000	100.0000	100.0000
region4	100.0000	0.0000	47.3684	0.0000
region5	100.0000	0.0000	100.0000	100.0000

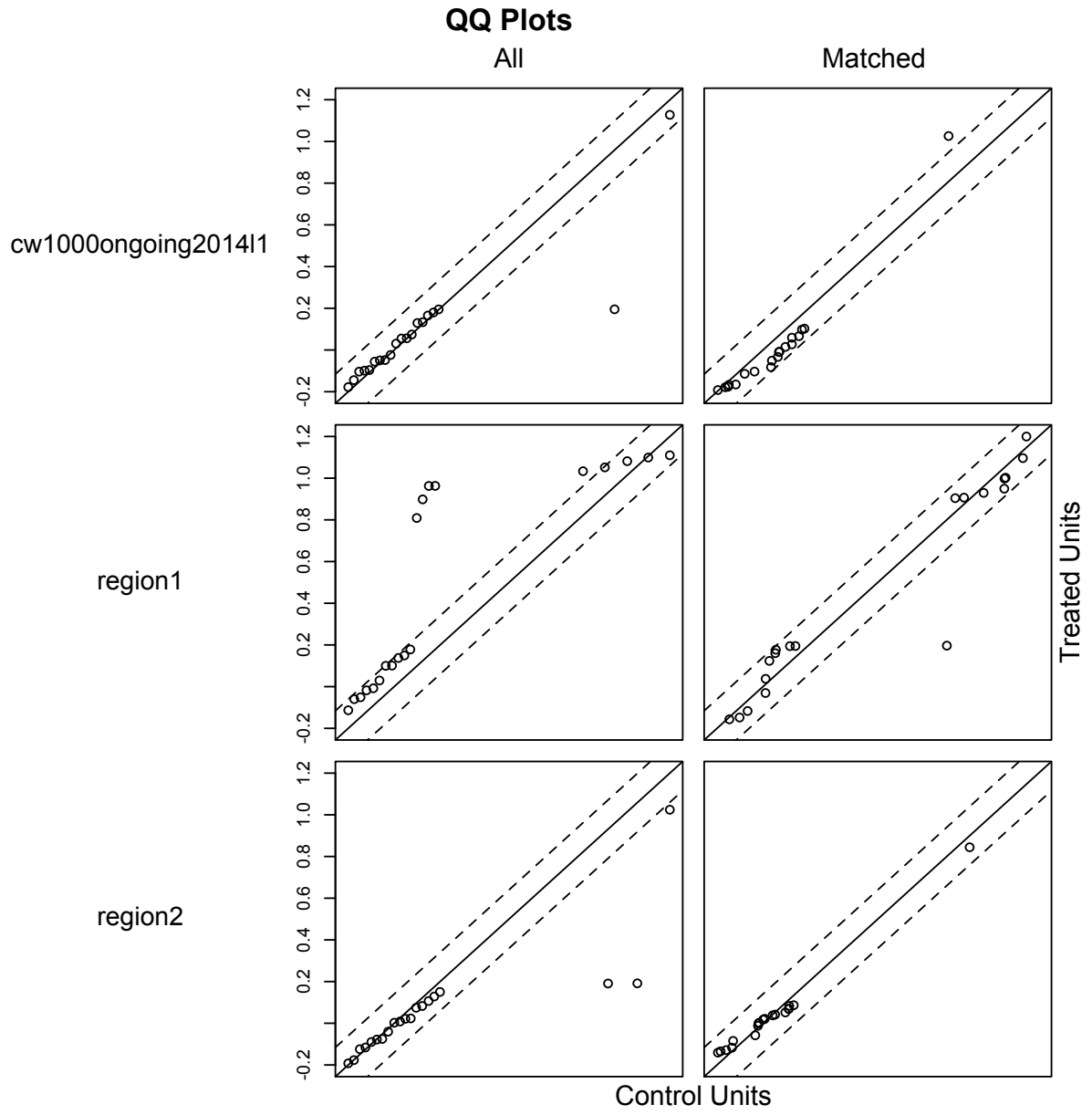
Sample sizes:

	Control	Treated
All	8318	20
Matched	19	20
Unmatched	8299	0
Discarded	0	0

QQ Plots





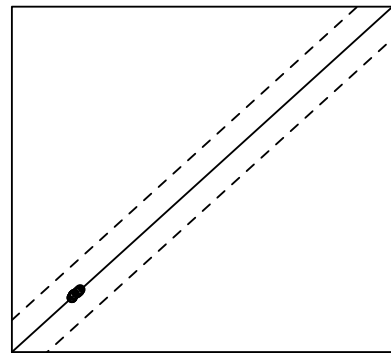
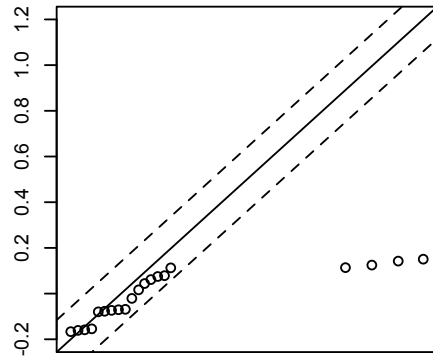


QQ Plots

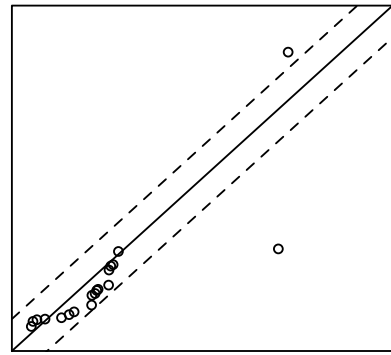
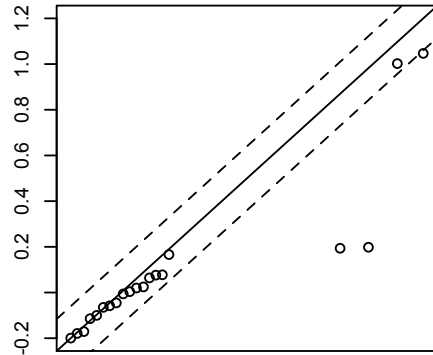
All

Matched

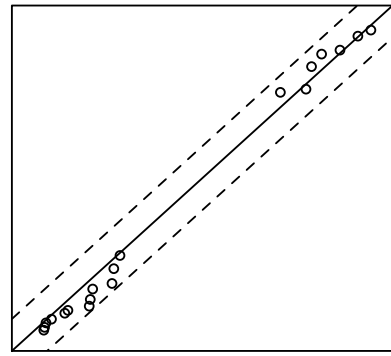
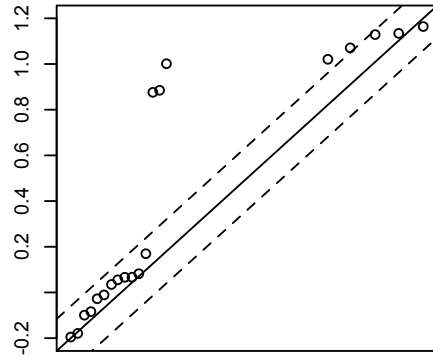
region3



region4



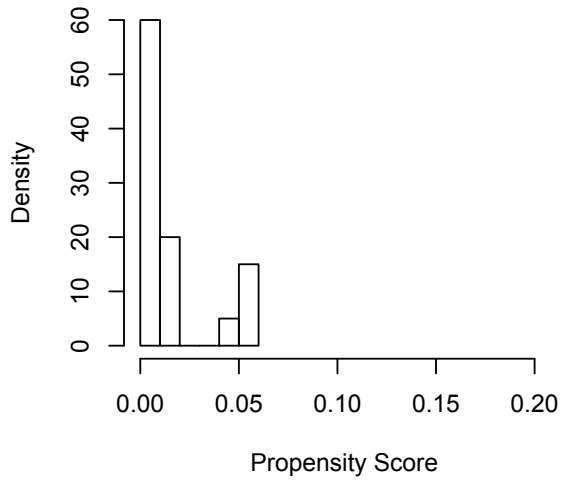
region5



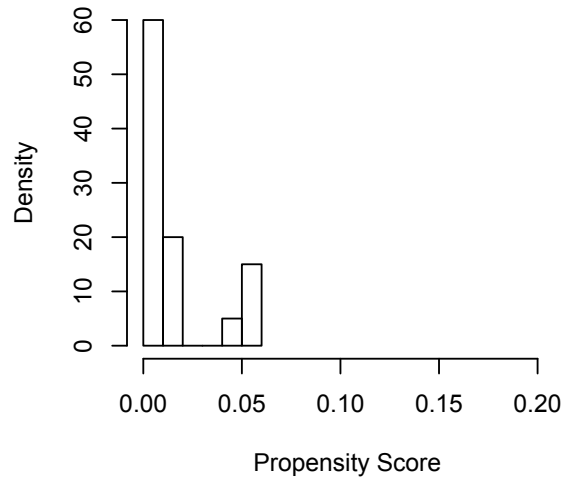
Treated Units

Control Units

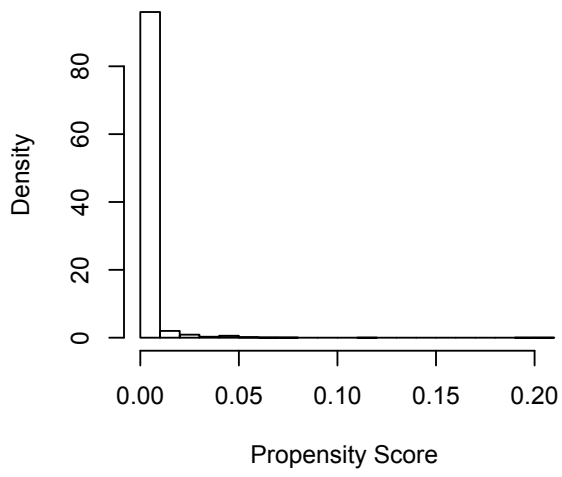
Raw Treated



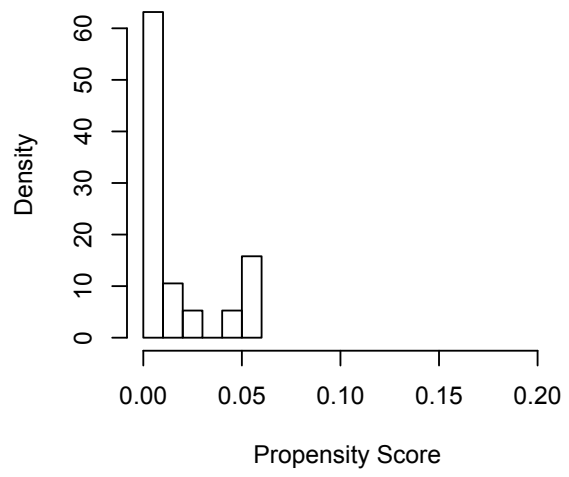
Matched Treated



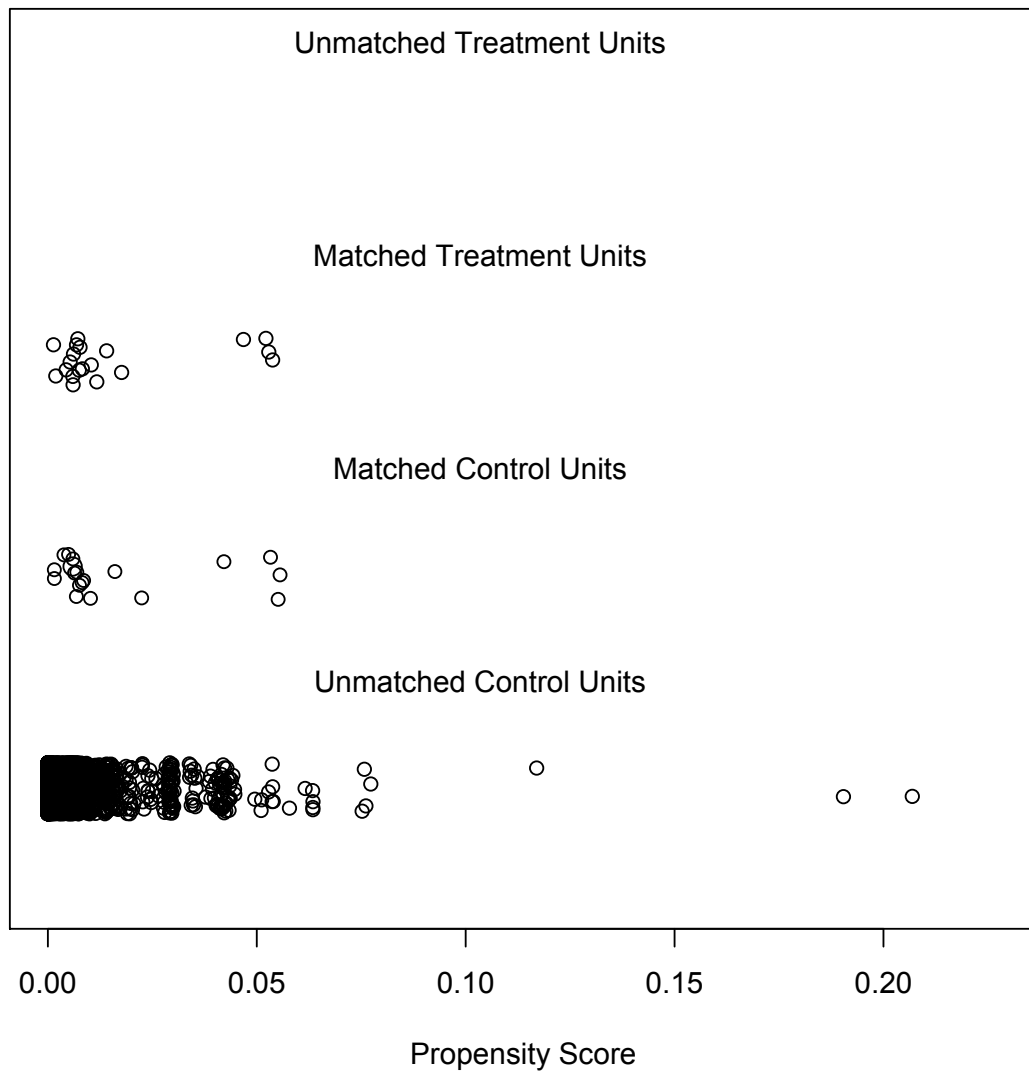
Raw Control



Matched Control



Distribution of Propensity Scores



INSTITUTIONAL FIRC: MATCHED ANALYSIS

tab civwar10_2014 abdstfirc, col chi2

civwar10_2014	abdstfirc		Total
	0	1	
0	19	16	35
	100.00	80.00	89.74
1	0	4	4
	0.00	20.00	10.26
Total	19	20	39
	100.00	100.00	100.00

Pearson chi2(1) = 4.2343 Pr = 0.040

tab civwar5_2014 abdstfirc, col chi2

civwar5_2014	abdstfirc		Total
	0	1	
0	19	16	35
	100.00	80.00	89.74
1	0	4	4
	0.00	20.00	10.26
Total	19	20	39
	100.00	100.00	100.00

Pearson chi2(1) = 4.2343 Pr = 0.040

ttest civwar10_2014, by(abdstfirc)

Two-sample t test with equal variances

Group	Obs	Mean	Std. Err.	Std. Dev.	[95% Conf. Interval]	
0	19	0	0	0	0	0
1	20	.2	.0917663	.4103913	.0079309	.3920691
combined	39	.1025641	.0492161	.3073547	.0029312	.202197
diff		-.2	.0942138		-.3908953	-.0091047

diff = mean(0) - mean(1) t = -2.1228
 Ho: diff = 0 degrees of freedom = 37

Ha: diff < 0 Pr(T < t) = 0.0203
 Ha: diff != 0 Pr(|T| > |t|) = 0.0405
 Ha: diff > 0 Pr(T > t) = 0.9797

ttest civwar5_2014, by(abdstfirc)

Two-sample t test with equal variances

Group	Obs	Mean	Std. Err.	Std. Dev.	[95% Conf. Interval]	
0	19	0	0	0	0	0
1	20	.2	.0917663	.4103913	.0079309	.3920691
combined	39	.1025641	.0492161	.3073547	.0029312	.202197
diff		-.2	.0942138		-.3908953	-.0091047

diff = mean(0) - mean(1) t = -2.1228
 Ho: diff = 0 degrees of freedom = 37

Ha: diff < 0 Pr(T < t) = 0.0203
 Ha: diff != 0 Pr(|T| > |t|) = 0.0405
 Ha: diff > 0 Pr(T > t) = 0.9797

INSTITUTIONAL FIRC (with interwar and gdbuffer)

Note: **peaceyrs** is omitted from matching because the balance between treated (34.15) and control (34.48) cases was already very good.

summary(instfirc.out3)

Call:

```
matchit(formula = abdstfirc ~ lnpec4_fill + lntpop4_banks +
  mountainous + newstate + democracy + interwar + cw1000ongoing201411 +
  gdbuffer + region1 + region2 + region3 + region4 + region5,
  data = instfirc, method = "genetic", pop.size = 150)
```

Summary of balance for all data:

	Means Treated	Means Control	SD Control	Mean Diff	eQQ Med	eQQ Mean	eQQ Max
distance	0.0210	0.0024	0.0073	0.0187	0.0076	0.0201	0.0941
lnpec4_fill	5.8619	7.0217	3.8684	-1.1598	0.7282	1.2457	4.4543
lntpop4_banks	8.1346	8.5581	1.8705	-0.4234	0.5903	0.7358	2.7548
mountainous	0.1615	0.1832	0.2210	-0.0216	0.0371	0.0704	0.5023
newstate	0.1000	0.0256	0.1580	0.0744	0.0000	0.0500	1.0000
democracy	0.0500	0.3557	0.4788	-0.3057	0.0000	0.3000	1.0000
interwar	0.3000	0.0513	0.2207	0.2487	0.0000	0.2500	1.0000
cw1000ongoing201411	0.0500	0.0908	0.2873	-0.0408	0.0000	0.0500	1.0000
gdbuffer	0.4000	0.1193	0.3241	0.2807	0.0000	0.2500	1.0000
region1	0.4500	0.2224	0.4159	0.2276	0.0000	0.2000	1.0000
region2	0.0500	0.1566	0.3635	-0.1066	0.0000	0.1000	1.0000
region3	0.0000	0.2046	0.4034	-0.2046	0.0000	0.2000	1.0000
region4	0.1000	0.1910	0.3931	-0.0910	0.0000	0.1000	1.0000
region5	0.4000	0.2253	0.4178	0.1747	0.0000	0.1500	1.0000

Summary of balance for matched data:

	Means Treated	Means Control	SD Control	Mean Diff	eQQ Med	eQQ Mean	eQQ Max
distance	0.0210	0.0212	0.0287	-0.0001	0.0004	0.0045	0.0528
lnpec4_fill	5.8619	5.5595	4.2897	0.3024	0.4592	0.7517	3.3398
lntpop4_banks	8.1346	8.0565	1.7940	0.0781	0.3077	0.3226	0.6971
mountainous	0.1615	0.1873	0.1755	-0.0258	0.0253	0.0460	0.1517
newstate	0.1000	0.1000	0.3082	0.0000	0.0000	0.0526	1.0000
democracy	0.0500	0.0500	0.2239	0.0000	0.0000	0.0000	0.0000
interwar	0.3000	0.3000	0.4708	0.0000	0.0000	0.0526	1.0000
cw1000ongoing201411	0.0500	0.0500	0.2239	0.0000	0.0000	0.0000	0.0000
gdbuffer	0.4000	0.4000	0.5033	0.0000	0.0000	0.0526	1.0000
region1	0.4500	0.4500	0.5111	0.0000	0.0000	0.0526	1.0000
region2	0.0500	0.0500	0.2239	0.0000	0.0000	0.0000	0.0000
region3	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
region4	0.1000	0.1000	0.3082	0.0000	0.0000	0.0526	1.0000
region5	0.4000	0.4000	0.5033	0.0000	0.0000	0.0000	0.0000

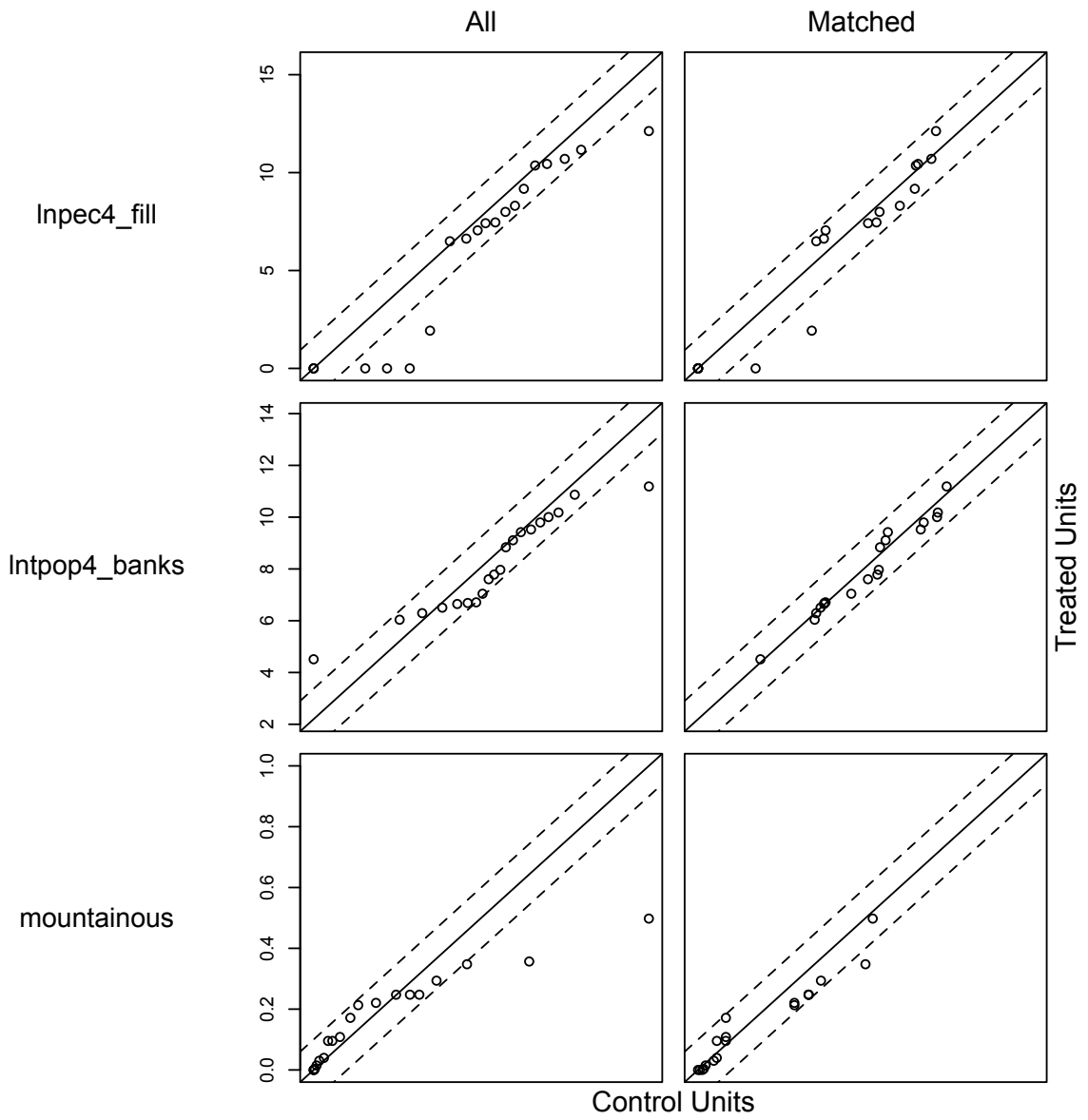
Percent Balance Improvement:

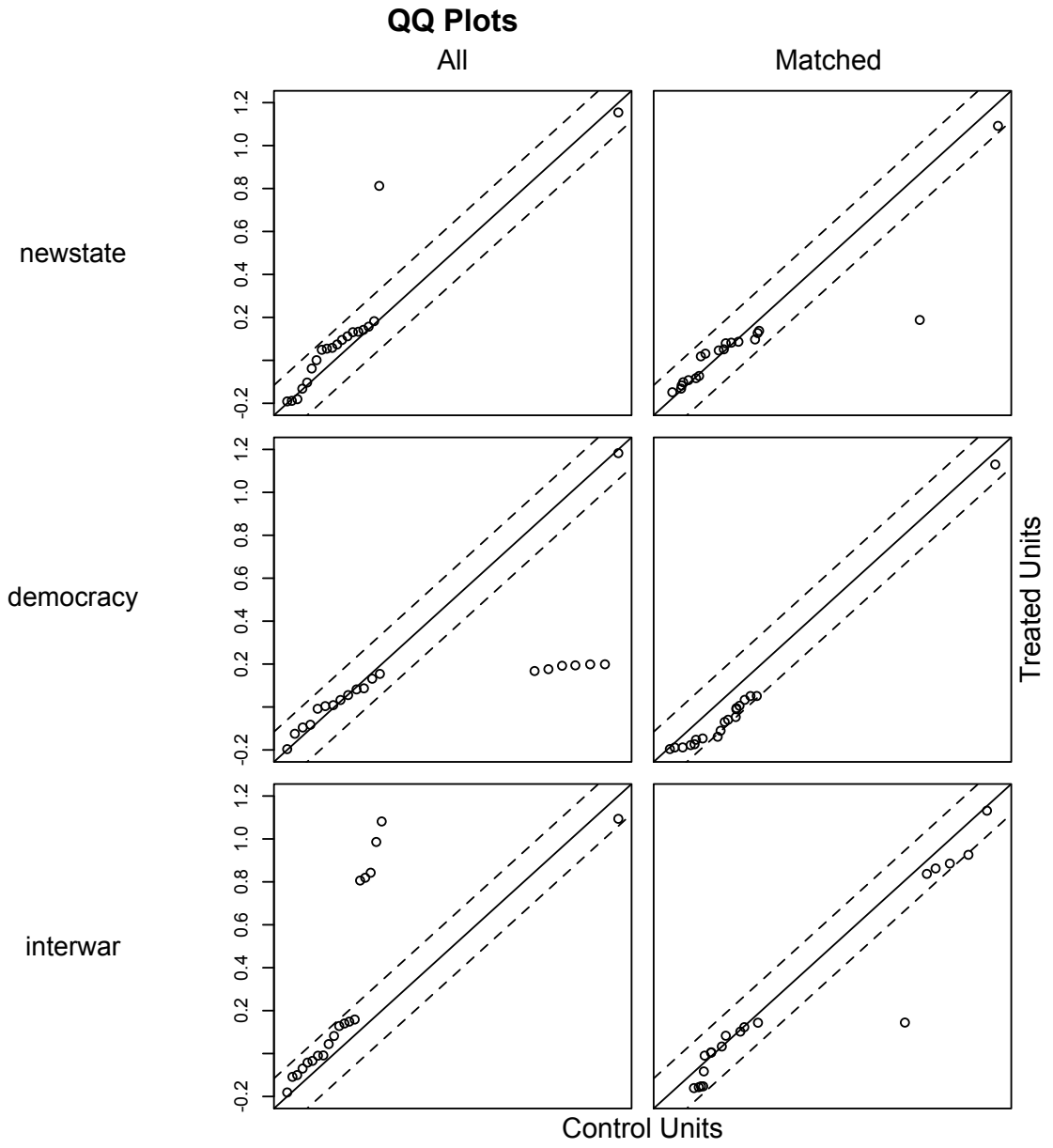
	Mean Diff.	eQQ Med	eQQ Mean	eQQ Max
distance	99.2969	94.9021	77.6965	43.8966
lnpec4_fill	73.9274	36.9428	39.6566	25.0215
lntpop4_banks	81.5453	47.8685	56.1599	74.6952
mountainous	-19.3395	31.6515	34.6687	69.7931
newstate	100.0000	0.0000	-5.2632	0.0000
democracy	100.0000	0.0000	100.0000	100.0000
interwar	100.0000	0.0000	78.9474	0.0000
cw1000ongoing201411	100.0000	0.0000	100.0000	100.0000
gdbuffer	100.0000	0.0000	78.9474	0.0000
region1	100.0000	0.0000	73.6842	0.0000
region2	100.0000	0.0000	100.0000	100.0000
region3	100.0000	0.0000	100.0000	100.0000
region4	100.0000	0.0000	47.3684	0.0000
region5	100.0000	0.0000	100.0000	100.0000

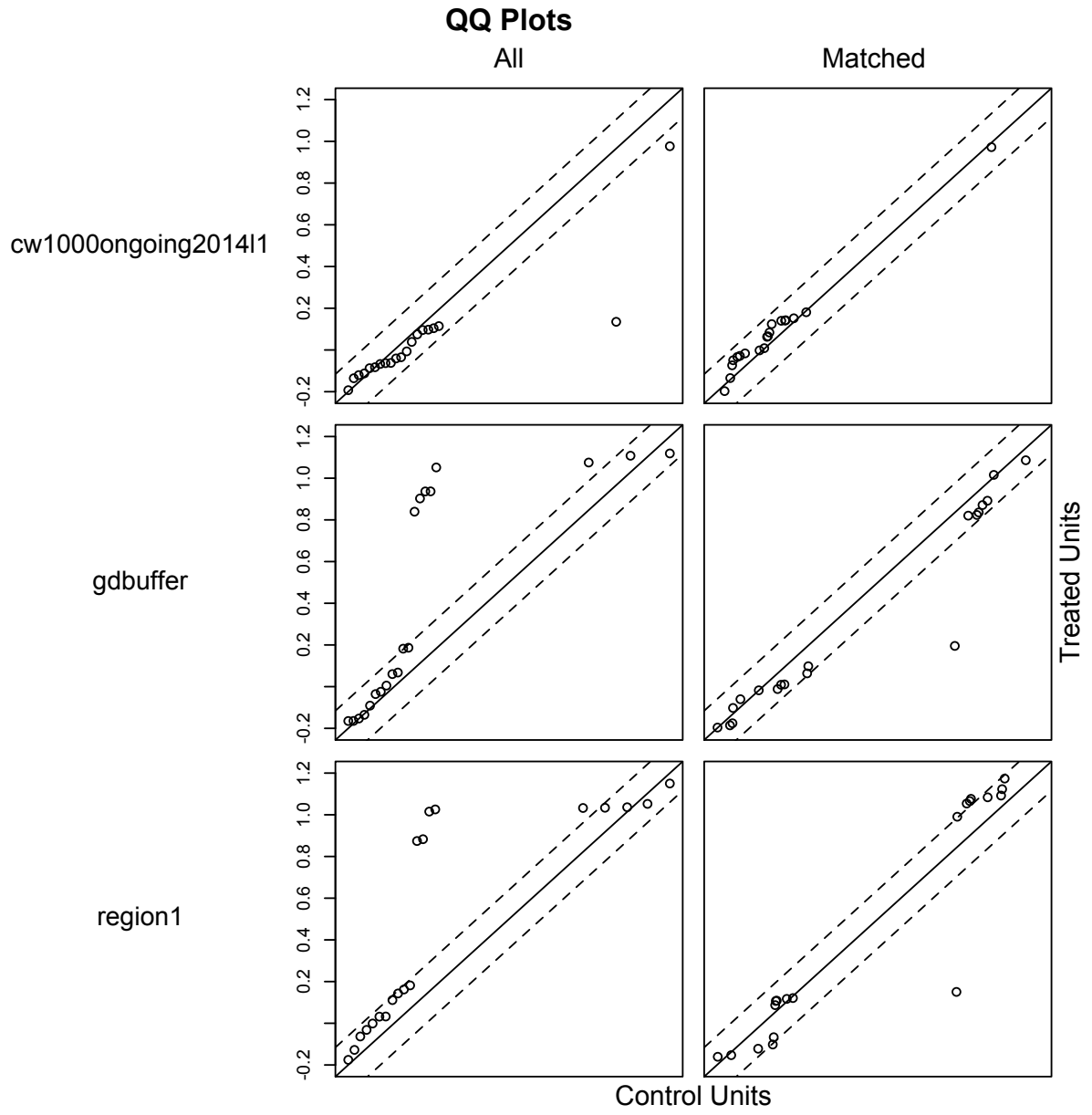
Sample sizes:

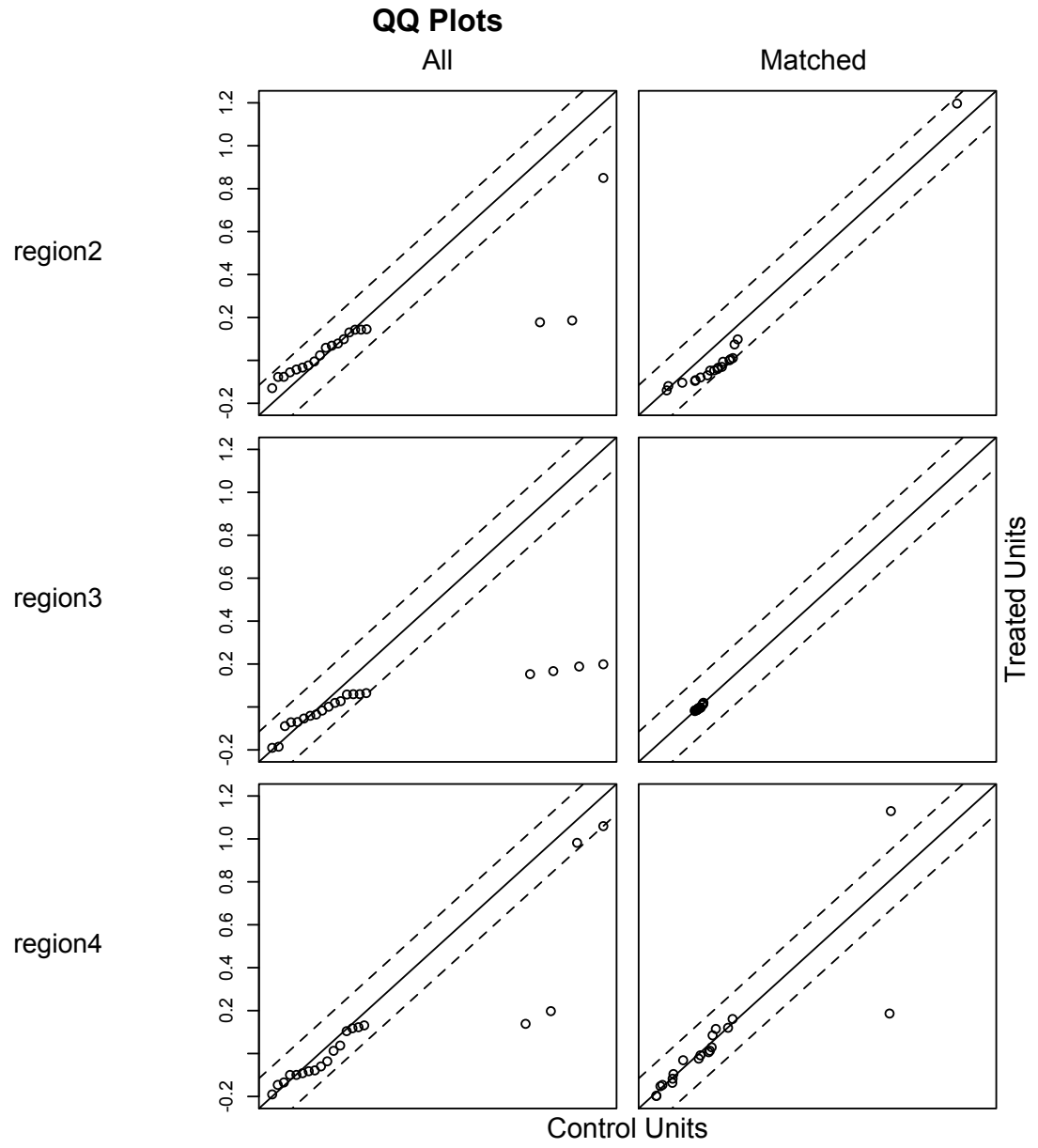
	Control	Treated
All	8318	20
Matched	19	20
Unmatched	8299	0
Discarded	0	0

QQ Plots

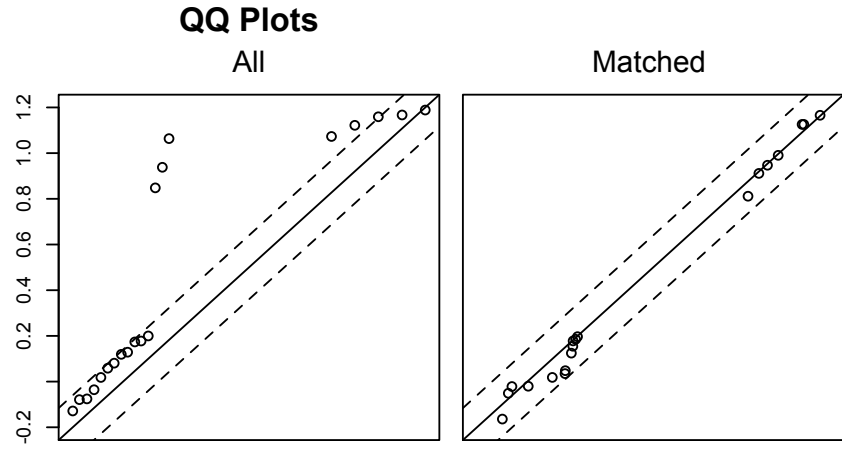








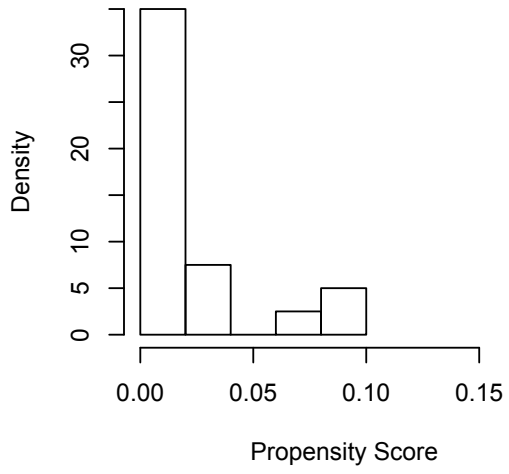
region5



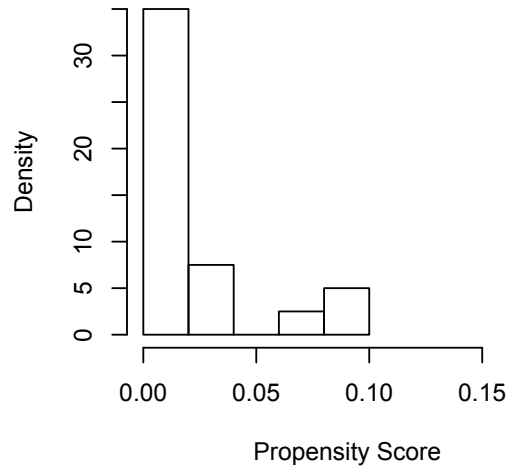
Treated Units

Control Units

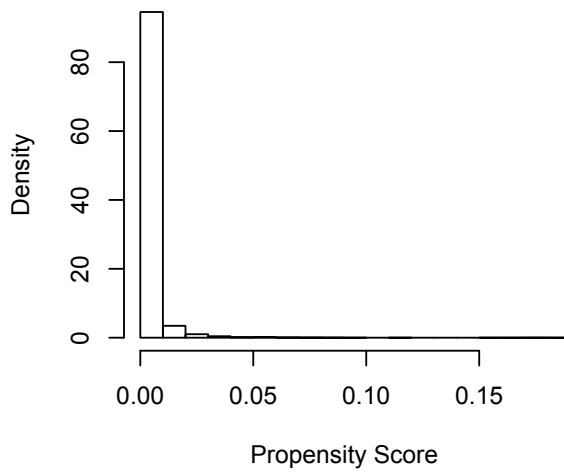
Raw Treated



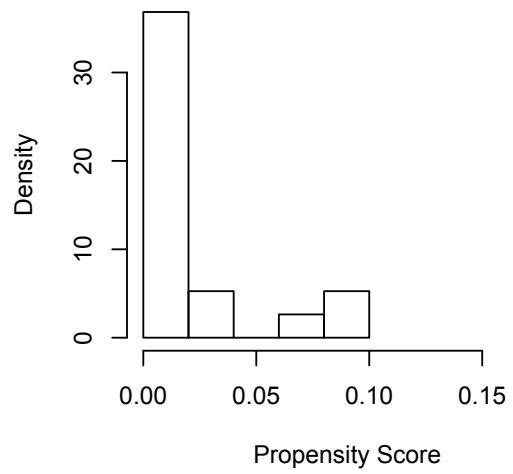
Matched Treated



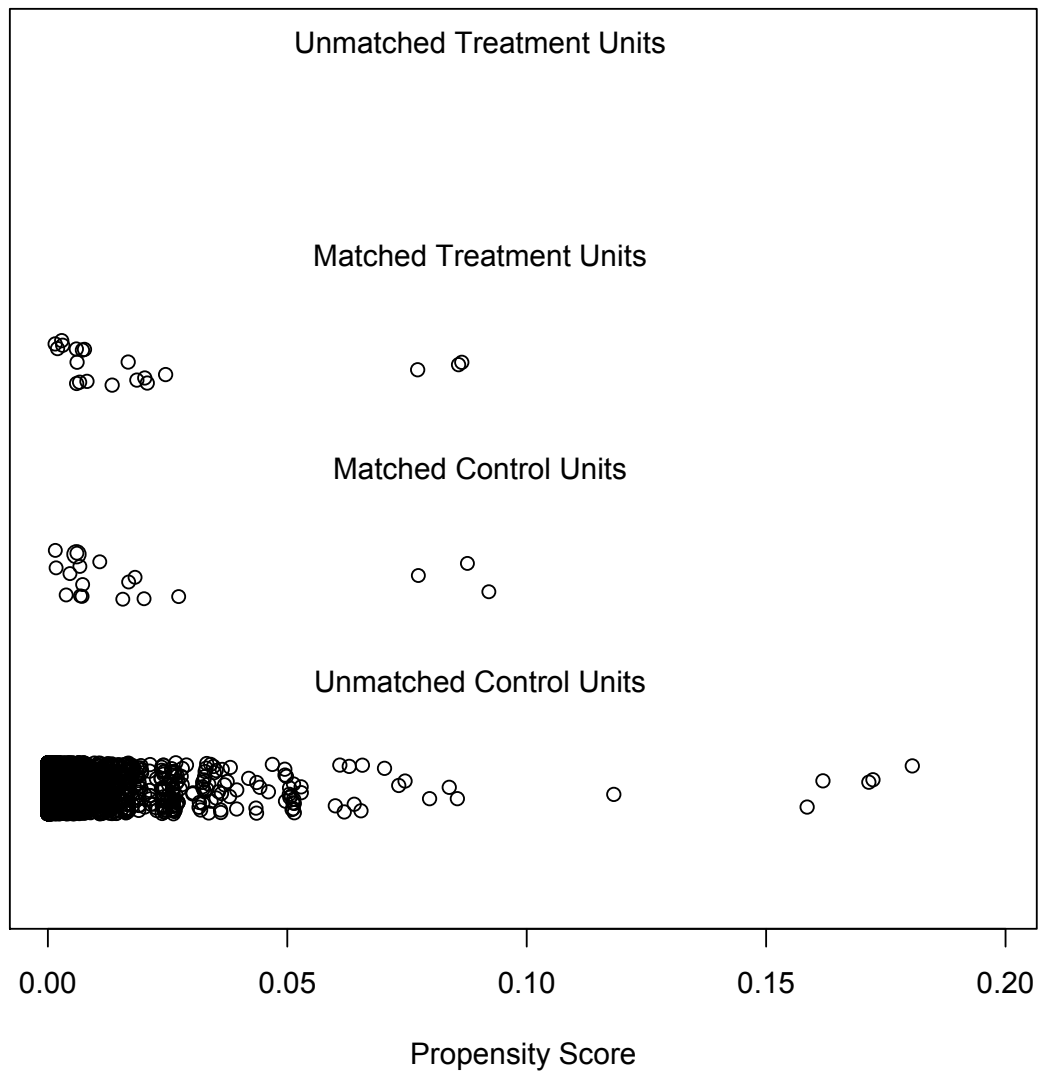
Raw Control



Matched Control



Distribution of Propensity Scores



INSTITUTIONAL FIRC: MATCHED ANALYSIS (with interwar and gdbuffer)

tab civwar10_2014 abdstfirc, col chi2

civwar10_2014	abdstfirc		Total
	0	1	
0	18 94.74	16 80.00	34 87.18
1	1 5.26	4 20.00	5 12.82
Total	19 100.00	20 100.00	39 100.00

Pearson chi2(1) = 1.8933 Pr = 0.169

tab civwar5_2014 abdstfirc, col chi2

civwar5_2014	abdstfirc		Total
	0	1	
0	18 94.74	16 80.00	34 87.18
1	1 5.26	4 20.00	5 12.82
Total	19 100.00	20 100.00	39 100.00

Pearson chi2(1) = 1.8933 Pr = 0.169

ttest civwar10_2014, by(abdstfirc)

Two-sample t test with equal variances

Group	Obs	Mean	Std. Err.	Std. Dev.	[95% Conf. Interval]	
0	19	.0526316	.0526316	.2294157	-.0579433	.1632064
1	20	.2	.0917663	.4103913	.0079309	.3920691
combined	39	.1282051	.0542336	.3386884	.018415	.2379952
diff		-.1473684	.1072571		-.3646919	.0699551
diff = mean(0) - mean(1)				t =	-1.3740	
Ho: diff = 0				degrees of freedom =	37	

Ha: diff < 0 Ha: diff != 0 Ha: diff > 0
 Pr(T < t) = 0.0889 Pr(|T| > |t|) = 0.1777 Pr(T > t) = 0.9111

ttest civwar5_2014, by(abdstfirc)

Two-sample t test with equal variances

Group	Obs	Mean	Std. Err.	Std. Dev.	[95% Conf. Interval]	
0	19	.0526316	.0526316	.2294157	-.0579433	.1632064
1	20	.2	.0917663	.4103913	.0079309	.3920691
combined	39	.1282051	.0542336	.3386884	.018415	.2379952
diff		-.1473684	.1072571		-.3646919	.0699551
diff = mean(0) - mean(1)				t =	-1.3740	
Ho: diff = 0				degrees of freedom =	37	

Ha: diff < 0 Ha: diff != 0 Ha: diff > 0
 Pr(T < t) = 0.0889 Pr(|T| > |t|) = 0.1777 Pr(T > t) = 0.9111

RESTORATION FIR

Note: **peaceyrs** (35.43 treated, 34.50 untreated), **mountainous** (.205 treated, .183 untreated), and **region3** (.214 treated, .205 untreated) were omitted because balance was already very good.

summary(restfirc.out3)

Call:

```
matchit(formula = abdstrestleadfirc ~ lnpec4_fill + lnpop4_banks +
  newstate + democracy + losewar10 + cw1000ongoing201411 +
  region1 + region2 + region4 + region5, data = restfirc, method = "genetic",
  pop.size = 150)
```

Summary of balance for all data:

	Means Treated	Means Control	SD Control	Mean Diff	eQQ Med	eQQ Mean	eQQ Max
distance	0.0782	0.0031	0.0101	0.0751	0.0365	0.0683	0.3130
lnpec4_fill	4.8666	7.0217	3.8684	-2.1551	1.8249	2.2043	5.1741
lnpop4_banks	8.0846	8.5581	1.8705	-0.4735	0.5175	0.8250	3.4588
newstate	0.0000	0.0256	0.1580	-0.0256	0.0000	0.0357	1.0000
democracy	0.2143	0.3557	0.4788	-0.1414	0.0000	0.1429	1.0000
losewar10	0.3929	0.0518	0.2217	0.3410	0.0000	0.3214	1.0000
cw1000ongoing201411	0.2143	0.0908	0.2873	0.1235	0.0000	0.1071	1.0000
region1	0.6786	0.2224	0.4159	0.4562	0.0000	0.4286	1.0000
region2	0.0714	0.1566	0.3635	-0.0852	0.0000	0.1071	1.0000
region4	0.0000	0.1910	0.3931	-0.1910	0.0000	0.2143	1.0000
region5	0.0357	0.2253	0.4178	-0.1896	0.0000	0.2143	1.0000

Summary of balance for matched data:

	Means Treated	Means Control	SD Control	Mean Diff	eQQ Med	eQQ Mean	eQQ Max
distance	0.0782	0.0700	0.0798	0.0082	0.0104	0.0235	0.1613
lnpec4_fill	4.8666	5.2972	3.7251	-0.4306	0.5132	1.4936	5.4596
lnpop4_banks	8.0846	8.2116	1.6938	-0.1270	0.3424	0.3942	1.5129
newstate	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
democracy	0.2143	0.2500	0.4432	-0.0357	0.0000	0.0909	1.0000
losewar10	0.3929	0.3929	0.4999	0.0000	0.0000	0.0455	1.0000
cw1000ongoing201411	0.2143	0.2500	0.4432	-0.0357	0.0000	0.0455	1.0000
region1	0.6786	0.7143	0.4624	-0.0357	0.0000	0.0455	1.0000
region2	0.0714	0.0714	0.2636	0.0000	0.0000	0.0455	1.0000
region4	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
region5	0.0357	0.0357	0.1899	0.0000	0.0000	0.0000	0.0000

Percent Balance Improvement:

	Mean Diff.	eQQ Med	eQQ Mean	eQQ Max
distance	89.1050	71.4067	65.5683	48.4885
lnpec4_fill	80.0177	71.8801	32.2404	-5.5177
lnpop4_banks	73.1802	33.8390	52.2210	56.2602
newstate	100.0000	0.0000	100.0000	100.0000
democracy	74.7511	0.0000	36.3636	0.0000
losewar10	100.0000	0.0000	85.8586	0.0000
cw1000ongoing201411	71.0859	0.0000	57.5758	0.0000
region1	92.1707	0.0000	89.3939	0.0000
region2	100.0000	0.0000	57.5758	0.0000
region4	100.0000	0.0000	100.0000	100.0000
region5	100.0000	0.0000	100.0000	100.0000

Sample sizes:

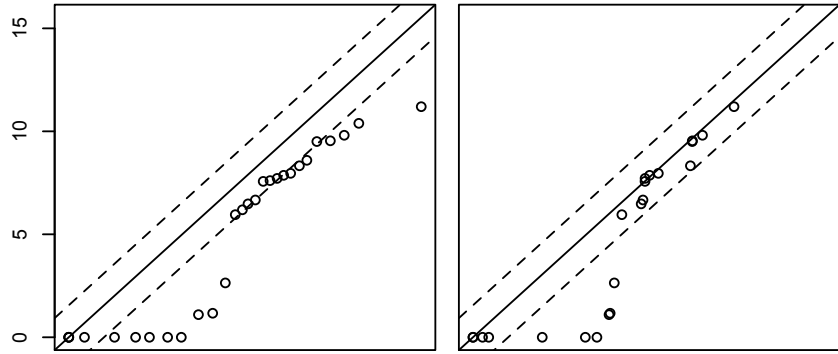
	Control	Treated
All	8318	28
Matched	22	28
Unmatched	8296	0
Discarded	0	0

QQ Plots

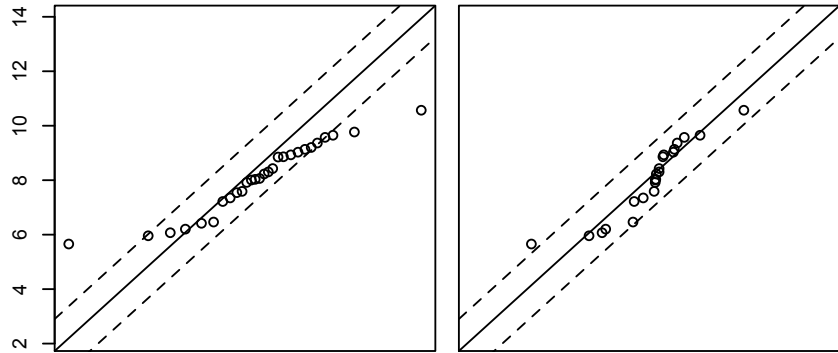
All

Matched

Inpec4_fill

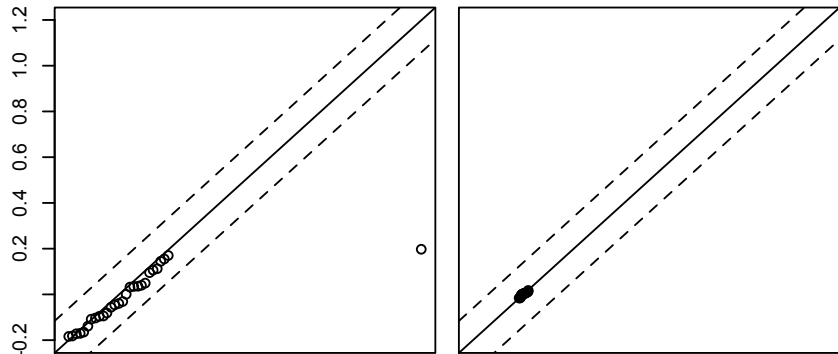


Intpop4_banks

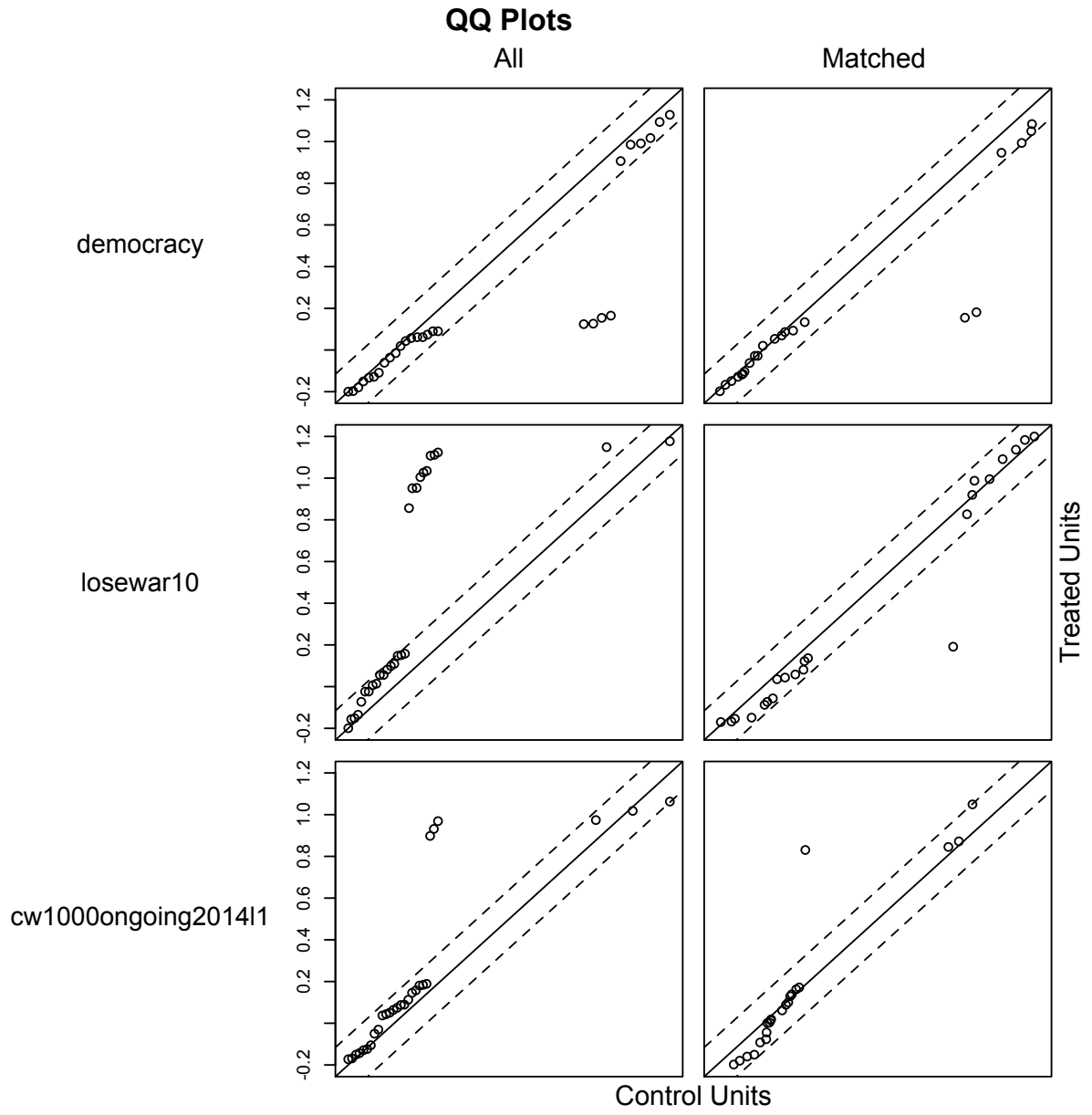


Treated Units

newstate



Control Units

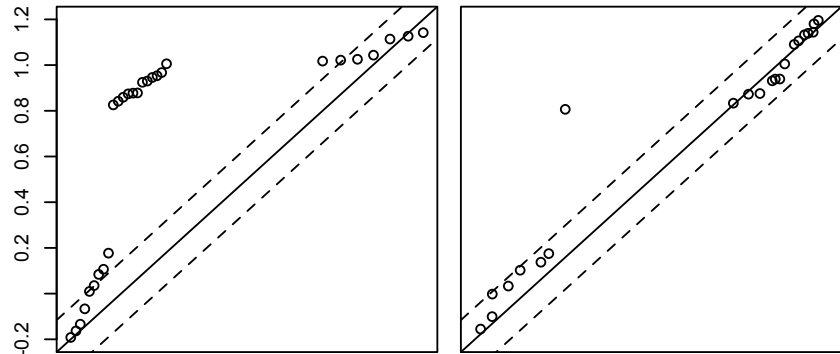


QQ Plots

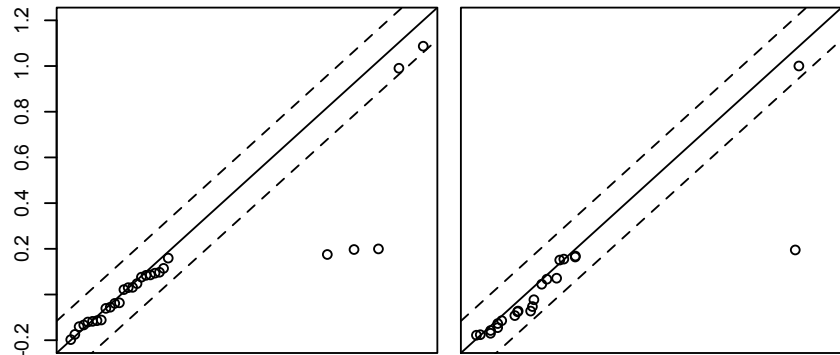
All

Matched

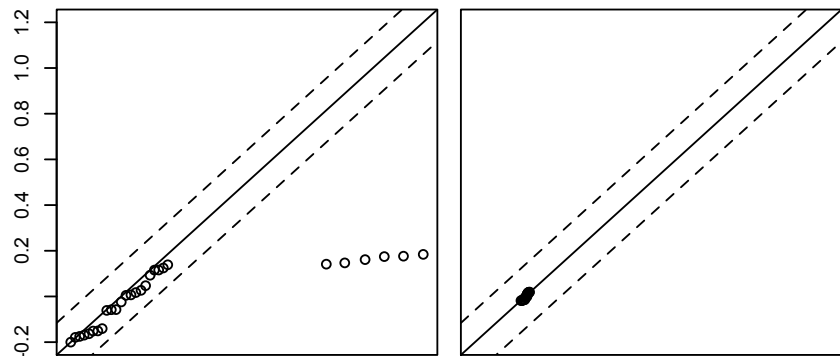
region1



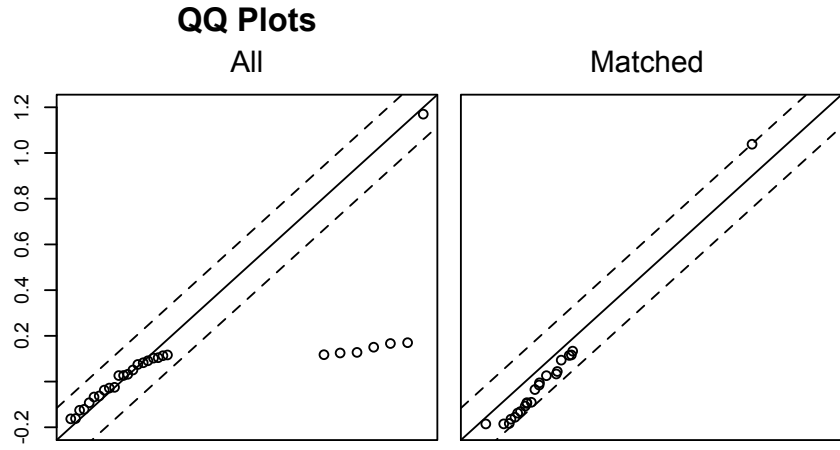
region2



region4



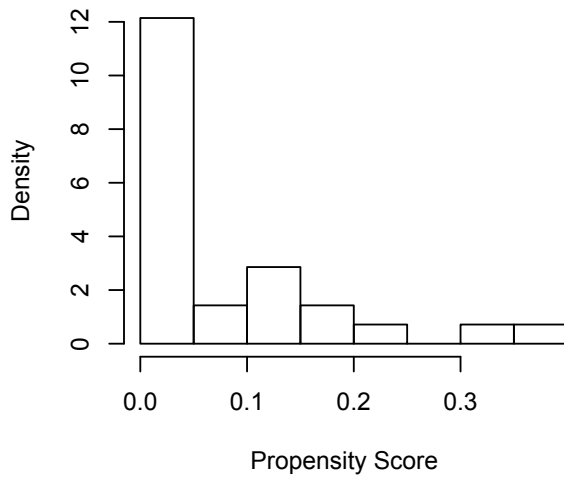
region5



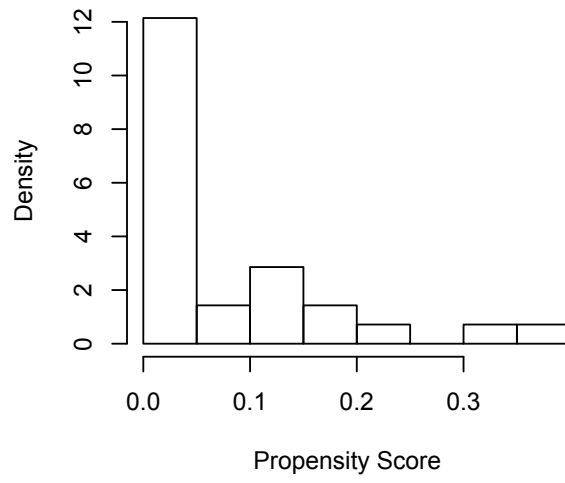
Treated Units

Control Units

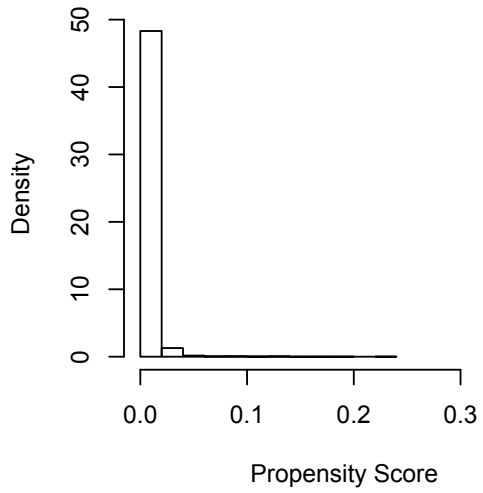
Raw Treated



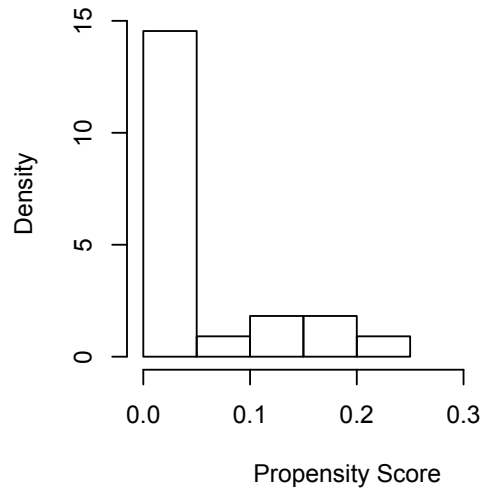
Matched Treated



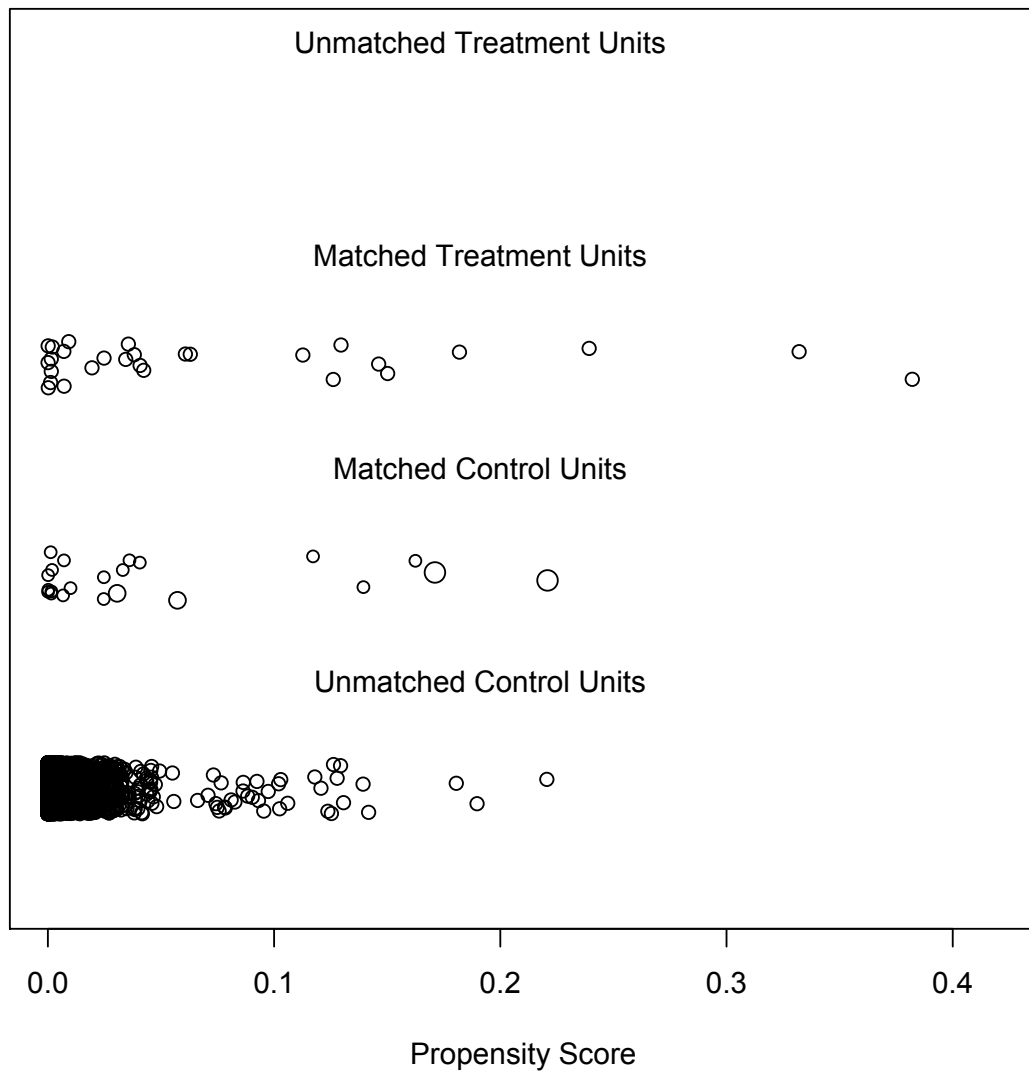
Raw Control



Matched Control



Distribution of Propensity Scores



RESTORATION FIRC: MATCHED ANALYSIS

tab civwar10_2014 abdrestdleadfirc, col chi2

civwar10_2014	abdrestdleadfirc		Total
	0	1	
0	20 90.91	25 89.29	45 90.00
1	2 9.09	3 10.71	5 10.00
Total	22 100.00	28 100.00	50 100.00

Pearson chi2(1) = 0.0361 Pr = 0.849

tab civwar5_2014 abdrestdleadfirc, col chi2

civwar5_2014	abdrestdleadfirc		Total
	0	1	
0	20 90.91	26 92.86	46 92.00
1	2 9.09	2 7.14	4 8.00
Total	22 100.00	28 100.00	50 100.00

Pearson chi2(1) = 0.0635 Pr = 0.801

ttest civwar10_2014, by(abdrestdleadfirc)

Two-sample t test with equal variances

Group	Obs	Mean	Std. Err.	Std. Dev.	[95% Conf. Interval]	
0	22	.0909091	.0627332	.2942449	-.0395518	.22137
1	28	.1071429	.0595238	.3149704	-.0149899	.2292756
combined	50	.1	.0428571	.3030458	.0138753	.1861247
diff		-.0162338	.0872014		-.191564	.1590964
diff = mean(0) - mean(1)					t =	-0.1862
Ho: diff = 0					degrees of freedom =	48

Ha: diff < 0 Ha: diff != 0 Ha: diff > 0
 Pr(T < t) = 0.4266 Pr(|T| > |t|) = 0.8531 Pr(T > t) = 0.5734

ttest civwar5_2014, by(abdrestdleadfirc)

Two-sample t test with equal variances

Group	Obs	Mean	Std. Err.	Std. Dev.	[95% Conf. Interval]	
0	22	.0909091	.0627332	.2942449	-.0395518	.22137
1	28	.0714286	.0495635	.2622653	-.0302673	.1731244
combined	50	.08	.0387562	.2740475	.0021166	.1578834
diff		.0194805	.0788355		-.1390289	.1779899
diff = mean(0) - mean(1)					t =	0.2471
Ho: diff = 0					degrees of freedom =	48

Ha: diff < 0 Ha: diff != 0 Ha: diff > 0
 Pr(T < t) = 0.5971 Pr(|T| > |t|) = 0.8059 Pr(T > t) = 0.4029

RESTORATION FIRIC (with interwar and gdbuffer)

Note: **peaceyrs** (35.43 treated, 34.50 untreated), **mountainous** (.205 treated, .183 untreated), and **region3** (.214 treated, .205 untreated) are omitted because the balance between treated and control cases was already very good.

summary(restfirc.out4)

Call:

```
matchit(formula = abdstrestleadfirc ~ lnpec4_fill + lntpop4_banks +
  newstate + democracy + interwar + cw1000ongoing201411 + gdbuffer +
  region1 + region2 + region4 + region5, data = restfirc, method = "genetic",
  pop.size = 150)
```

Summary of balance for all data:

	Means Treated	Means Control	SD Control	Mean Diff	eQQ Med	eQQ Mean	eQQ Max
distance	0.0527	0.0032	0.0084	0.0496	0.0204	0.0436	0.2570
lnpec4_fill	4.8666	7.0217	3.8684	-2.1551	1.8249	2.2043	5.1741
lntpop4_banks	8.0846	8.5581	1.8705	-0.4735	0.5175	0.8250	3.4588
newstate	0.0000	0.0256	0.1580	-0.0256	0.0000	0.0357	1.0000
democracy	0.2143	0.3557	0.4788	-0.1414	0.0000	0.1429	1.0000
interwar	0.2143	0.0513	0.2207	0.1630	0.0000	0.1429	1.0000
cw1000ongoing201411	0.2143	0.0908	0.2873	0.1235	0.0000	0.1071	1.0000
gdbuffer	0.3571	0.1193	0.3241	0.2379	0.0000	0.2143	1.0000
region1	0.6786	0.2224	0.4159	0.4562	0.0000	0.4286	1.0000
region2	0.0714	0.1566	0.3635	-0.0852	0.0000	0.1071	1.0000
region4	0.0000	0.1910	0.3931	-0.1910	0.0000	0.2143	1.0000
region5	0.0357	0.2253	0.4178	-0.1896	0.0000	0.2143	1.0000

Summary of balance for matched data:

	Means Treated	Means Control	SD Control	Mean Diff	eQQ Med	eQQ Mean	eQQ Max
distance	0.0527	0.0431	0.0566	0.0097	0.0023	0.0182	0.2570
lnpec4_fill	4.8666	5.0744	3.8734	-0.2078	0.3813	1.2669	5.9081
lntpop4_banks	8.0846	8.1601	1.9082	-0.0755	0.4935	0.5982	1.5129
newstate	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
democracy	0.2143	0.2500	0.4423	-0.0357	0.0000	0.0000	0.0000
interwar	0.2143	0.1786	0.3912	0.0357	0.0000	0.0000	0.0000
cw1000ongoing201411	0.2143	0.2500	0.4423	-0.0357	0.0000	0.0000	0.0000
gdbuffer	0.3571	0.3929	0.4989	-0.0357	0.0000	0.1250	1.0000
region1	0.6786	0.6786	0.4771	0.0000	0.0000	0.0417	1.0000
region2	0.0714	0.0714	0.2631	0.0000	0.0000	0.0417	1.0000
region4	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
region5	0.0357	0.0357	0.1896	0.0000	0.0000	0.0000	0.0000

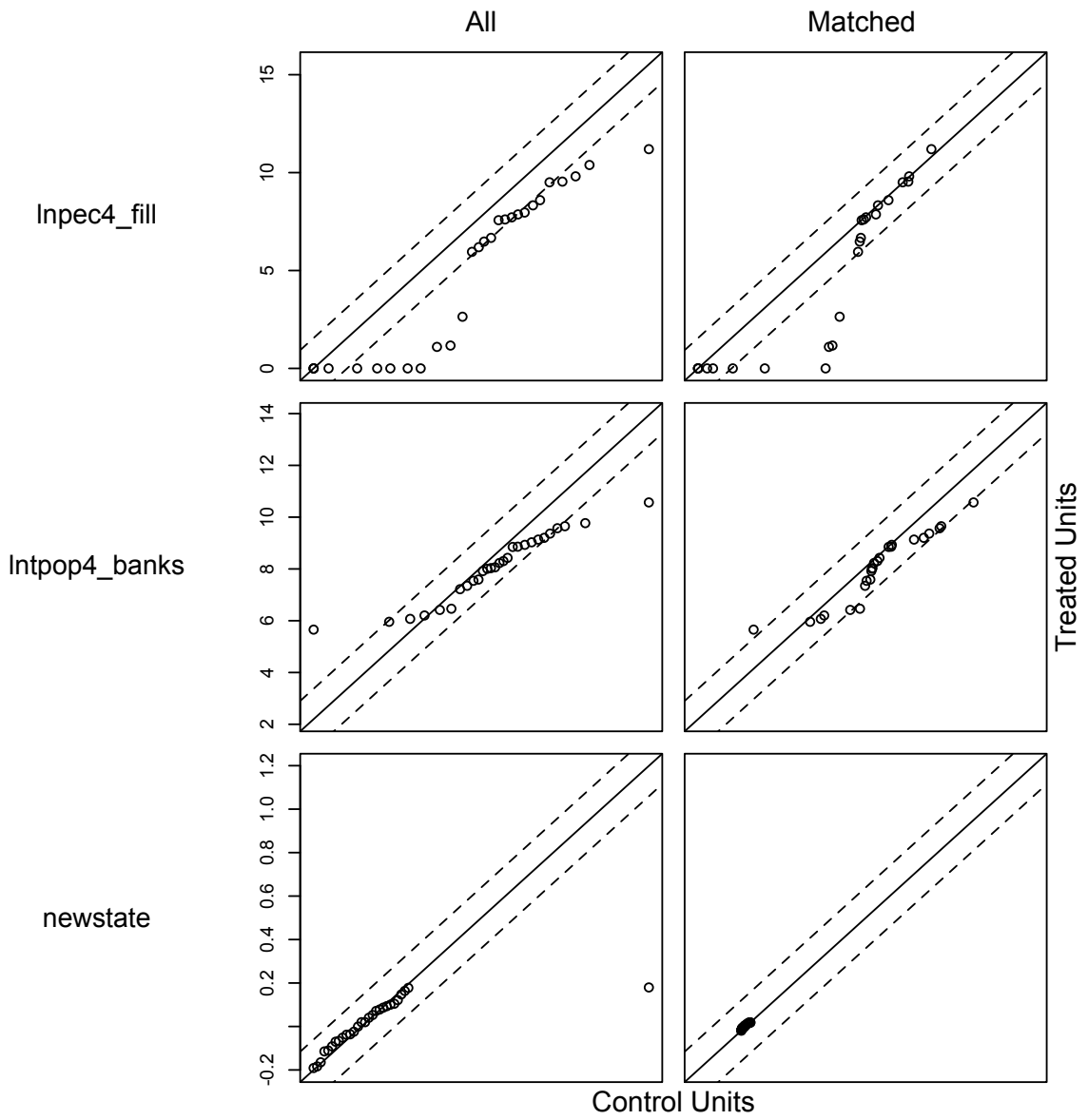
Percent Balance Improvement:

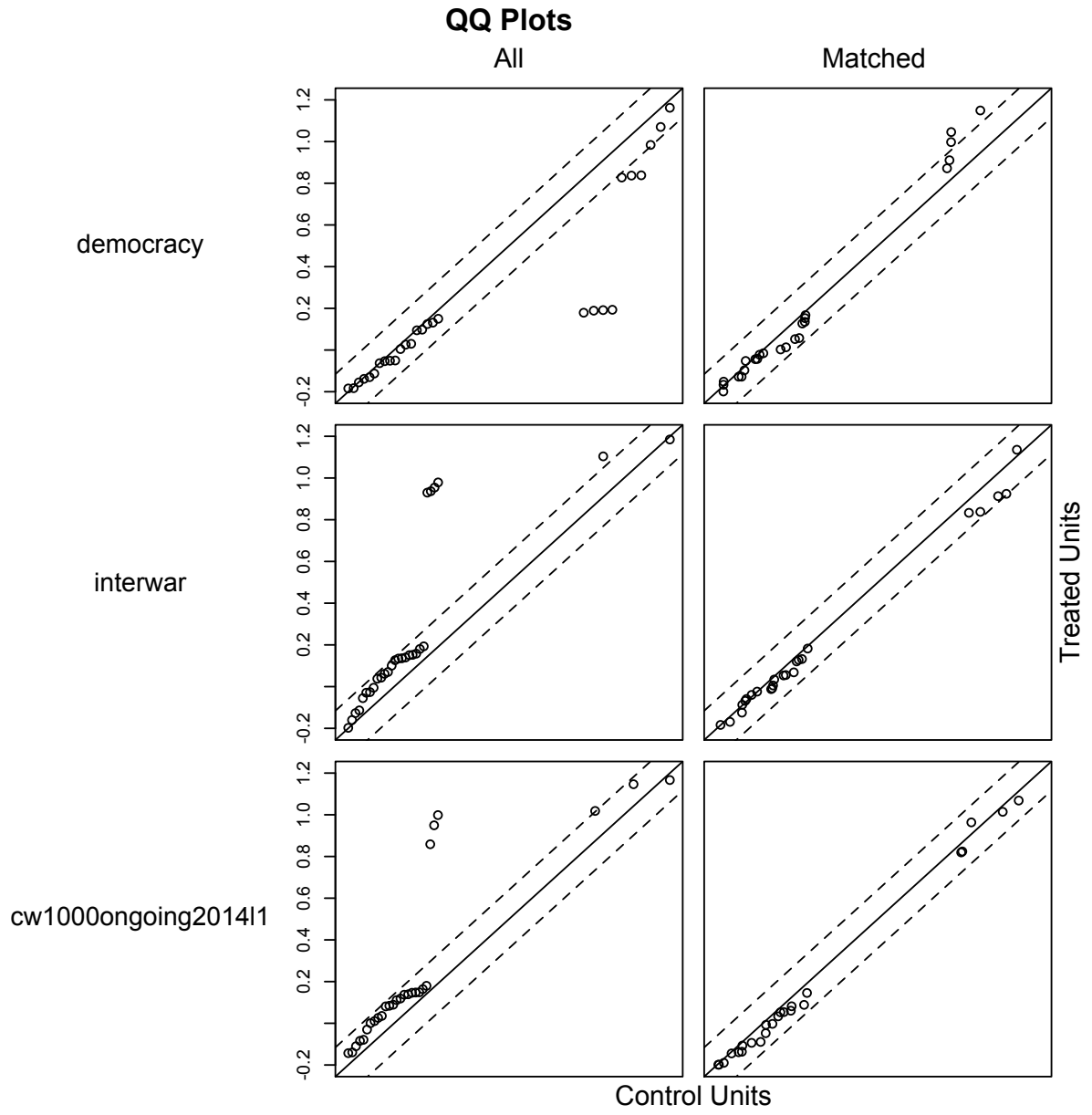
	Mean Diff.	eQQ Med	eQQ Mean	eQQ Max
distance	80.4337	88.9282	58.2605	0.0000
lnpec4_fill	90.3574	79.1081	42.5262	-14.1858
lntpop4_banks	84.0454	4.6419	27.4931	56.2602
newstate	100.0000	0.0000	100.0000	100.0000
democracy	74.7511	0.0000	100.0000	100.0000
interwar	78.0828	0.0000	100.0000	100.0000
cw1000ongoing201411	71.0859	0.0000	100.0000	100.0000
gdbuffer	84.9866	0.0000	41.6667	0.0000
region1	100.0000	0.0000	90.2778	0.0000
region2	100.0000	0.0000	61.1111	0.0000
region4	100.0000	0.0000	100.0000	100.0000
region5	100.0000	0.0000	100.0000	100.0000

Sample sizes:

	Control	Treated
All	8318	28
Matched	24	28
Unmatched	8294	0
Discarded	0	0

QQ Plots



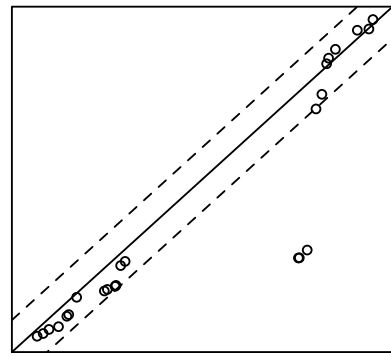
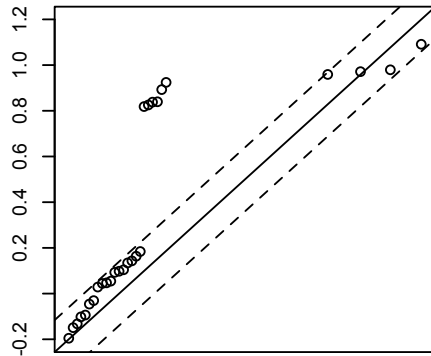


QQ Plots

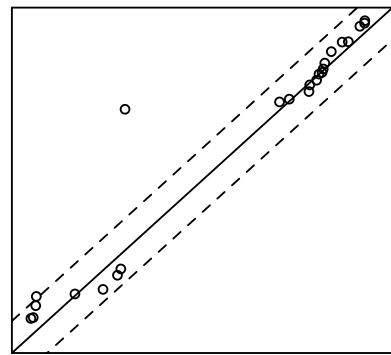
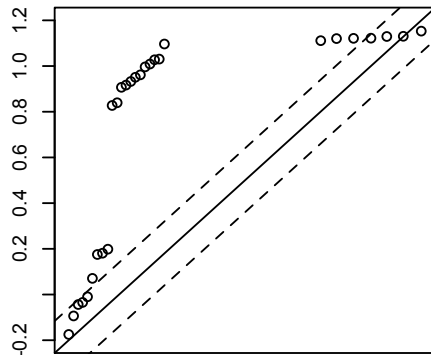
All

Matched

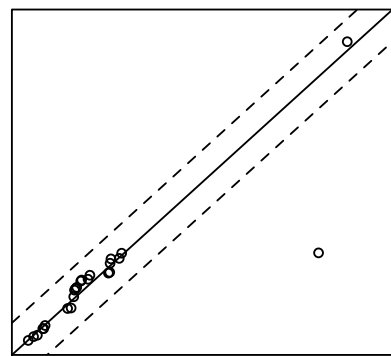
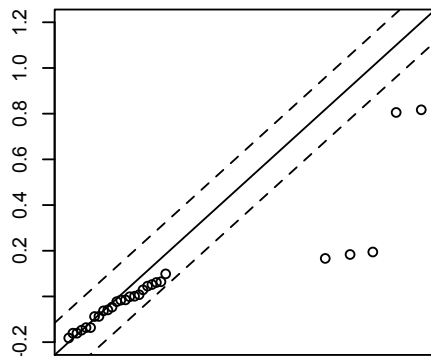
gdbuffer



region1



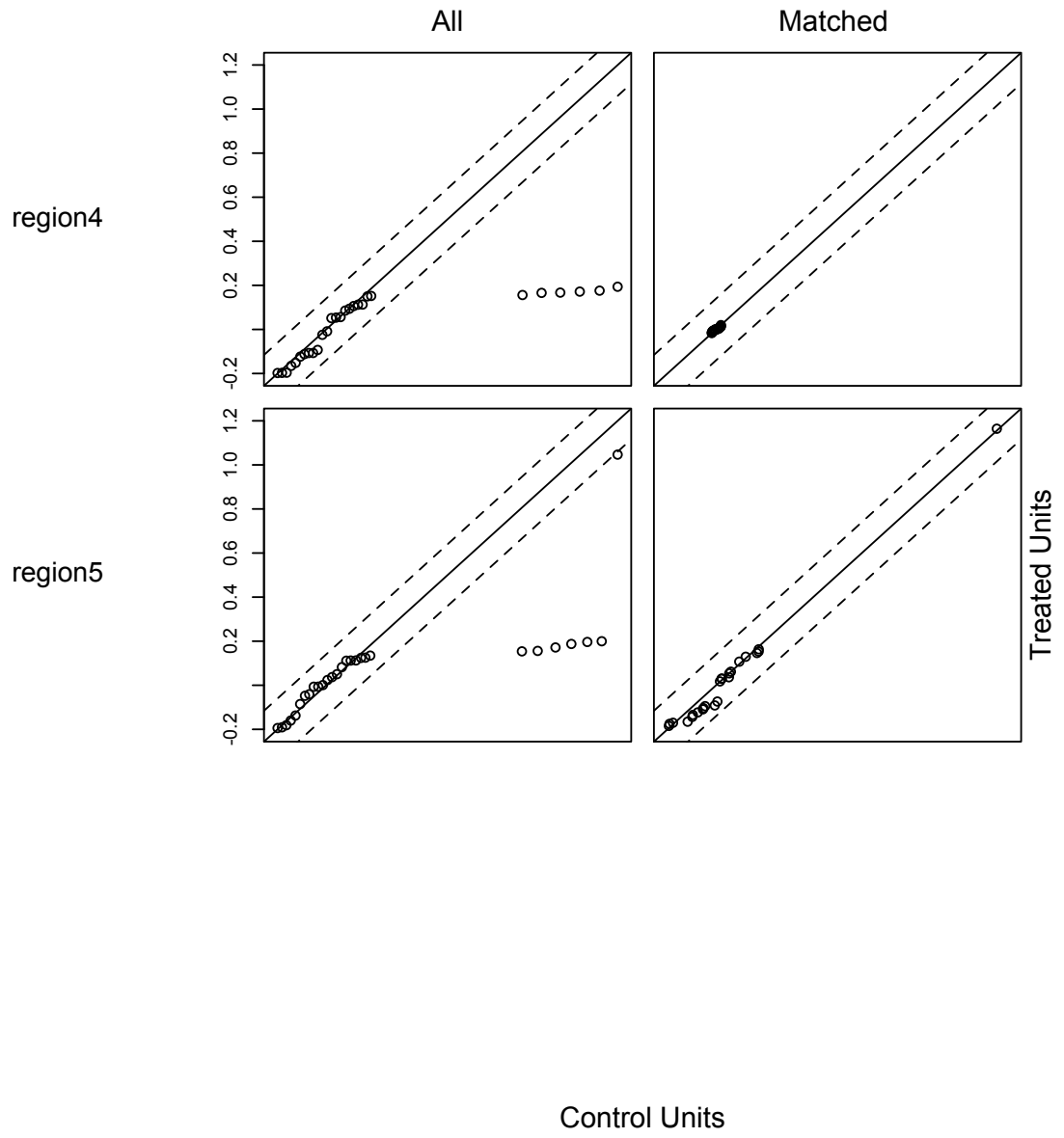
region2



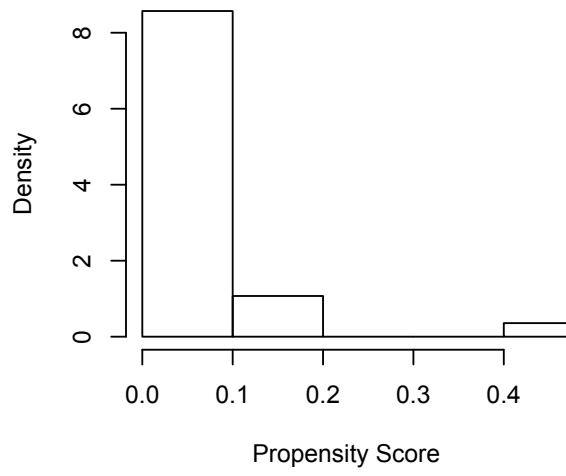
Treated Units

Control Units

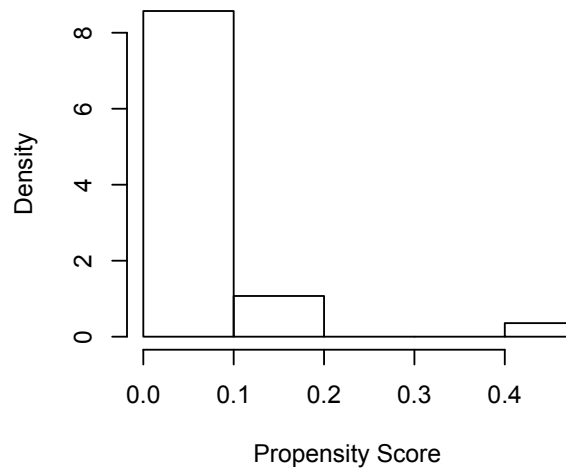
QQ Plots



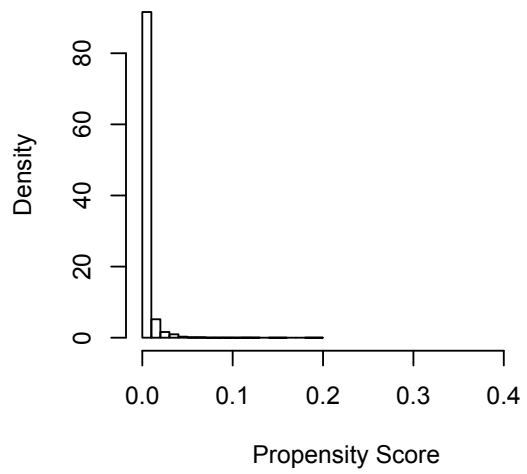
Raw Treated



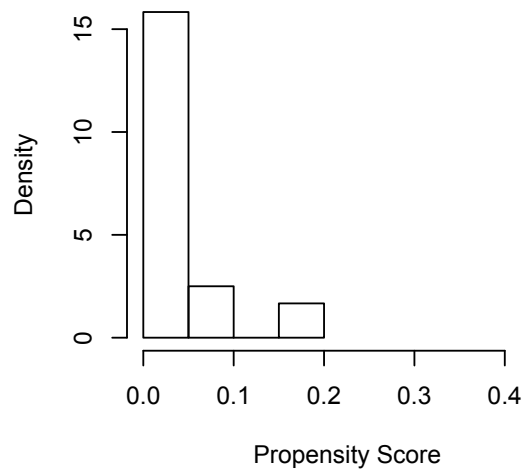
Matched Treated



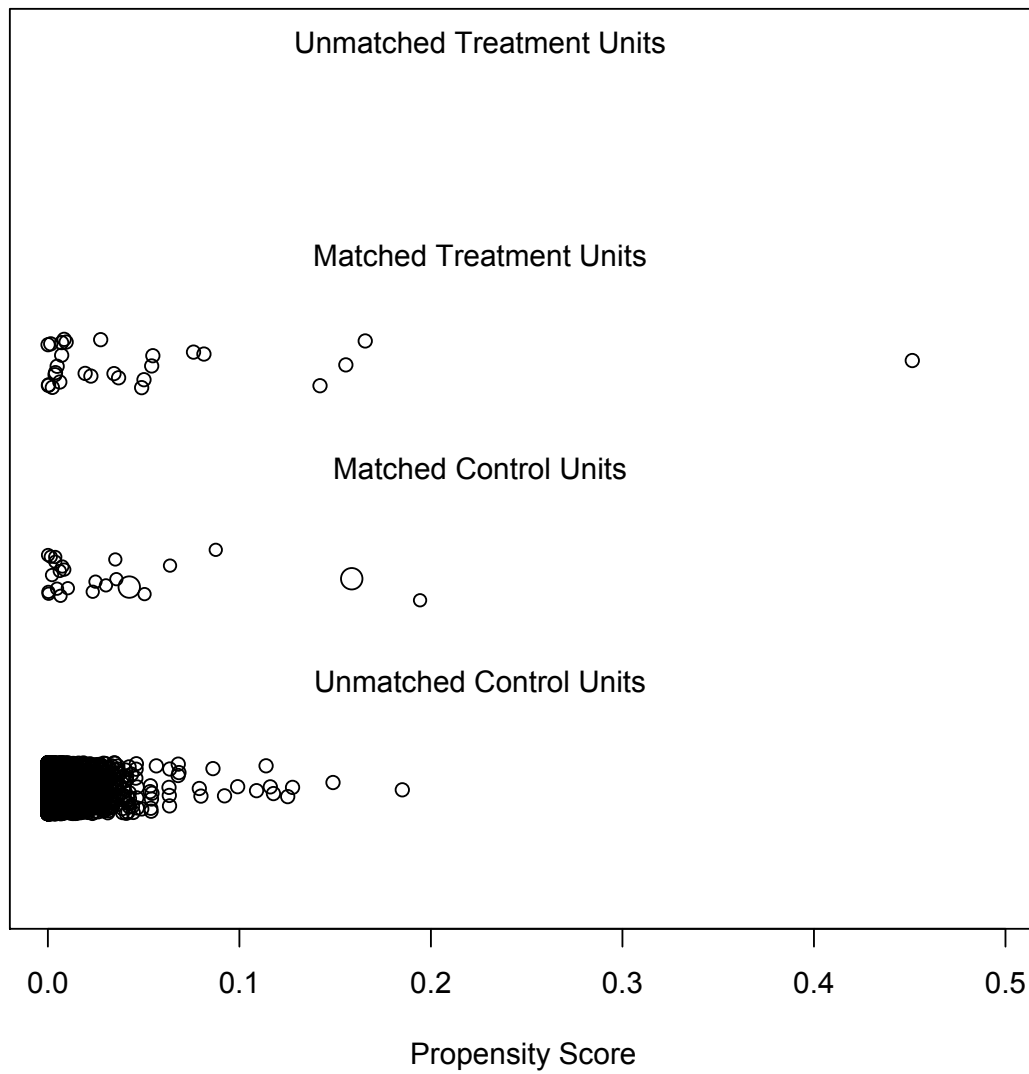
Raw Control



Matched Control



Distribution of Propensity Scores



RESTORATION FIRIC, MATCHED ANALYSIS (with interwar and gdbuffer)

tab civwar10_2014 abdrestdleadfirc, col chi2

civwar10_2014	abdrestdleadfirc		Total
	0	1	
0	20 83.33	25 89.29	45 86.54
1	4 16.67	3 10.71	7 13.46
Total	24 100.00	28 100.00	52 100.00

Pearson chi2(1) = 0.3930 Pr = 0.531

tab civwar5_2014 abdrestdleadfirc, col chi2

civwar5_2014	abdrestdleadfirc		Total
	0	1	
0	21 87.50	26 92.86	47 90.38
1	3 12.50	2 7.14	5 9.62
Total	24 100.00	28 100.00	52 100.00

Pearson chi2(1) = 0.4267 Pr = 0.514

ttest civwar10_2014, by(abdrestdleadfirc)

Two-sample t test with equal variances

Group	Obs	Mean	Std. Err.	Std. Dev.	[95% Conf. Interval]	
0	24	.1666667	.0777087	.3806935	.0059139	.3274194
1	28	.1071429	.0595238	.3149704	-.0149899	.2292756
combined	52	.1346154	.0477933	.3446423	.0386664	.2305644
diff		.0595238	.096458		-.1342177	.2532653

diff = mean(0) - mean(1) t = 0.6171
Ho: diff = 0 degrees of freedom = 50

Ha: diff < 0 Pr(T < t) = 0.7300
Ha: diff != 0 Pr(|T| > |t|) = 0.5400
Ha: diff > 0 Pr(T > t) = 0.2700

ttest civwar5_2014, by(abdrestdleadfirc)

Two-sample t test with equal variances

Group	Obs	Mean	Std. Err.	Std. Dev.	[95% Conf. Interval]	
0	24	.125	.0689597	.337832	-.0176539	.2676539
1	28	.0714286	.0495635	.2622653	-.0302673	.1731244
combined	52	.0961538	.0412806	.2976783	.0132797	.179028
diff		.0535714	.0832866		-.1137145	.2208574

diff = mean(0) - mean(1) t = 0.6432
Ho: diff = 0 degrees of freedom = 50

Ha: diff < 0 Pr(T < t) = 0.7385
Ha: diff != 0 Pr(|T| > |t|) = 0.5230
Ha: diff > 0 Pr(T > t) = 0.2615

