

**Summary of Balance for All Data:**

	<i>Means</i> <i>Treated</i>	<i>Means</i> <i>Control</i>	<i>Std. Mean</i> <i>Diff.</i>	<i>Var. Ratio</i>	<i>eCDF</i> <i>Mean</i>	<i>eCDF</i> <i>Max</i>
distance	0.544	0.276	1.077	1.62	0.301	0.462
i.age	40.51	39.628	0.096	1.141	0.037	0.109
i.sex	0.769	0.657	0.266	.	0.112	0.112
i.educ	4.548	4.983	-0.406	1.516	0.072	0.252
i.insu	1.481	1.477	0.004	1.262	0.017	0.041
i.drug	0.99	0.977	0.14	.	0.014	0.014
i.seve	2.76	2.529	0.263	0.858	0.058	0.09
com.t	2.933	1.715	0.81	1.526	0.135	0.381
pcs.sd	41.311	48.274	-0.665	1.309	0.204	0.377
mcs.sd	48.852	48.429	0.043	0.99	0.02	0.059

**Summary of Balance for Matched Data:**

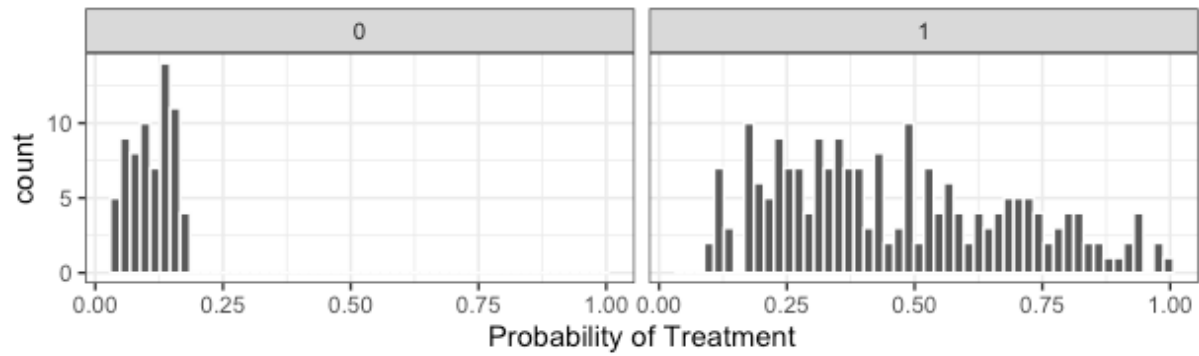
	<i>Means</i> <i>Treated</i>	<i>Means</i> <i>Control</i>	<i>Std. Mean</i> <i>Diff.</i>	<i>Var.</i> <i>Ratio</i>	<i>eCDF</i> <i>Mean</i>	<i>eCDF</i> <i>Max</i>	<i>Pair</i> <i>Dist.</i>
distance	0.544	0.384	0.641	1.913	0.142	0.375	0.642
i.age	40.51	40.635	-0.014	1.224	0.03	0.087	1.114
i.sex	0.769	0.731	0.091	.	0.038	0.038	0.913
i.educ	4.548	4.779	-0.216	1.371	0.042	0.163	0.917
i.insu	1.481	1.346	0.138	2.756	0.026	0.048	0.651
i.drug	0.99	1	-0.099	.	0.01	0.01	0.099
i.seve	2.76	2.615	0.165	0.801	0.036	0.077	1.197
com.t	2.933	2.163	0.512	1.399	0.085	0.25	0.717
pcs.sd	41.311	45.471	-0.397	1.254	0.116	0.24	0.929
mcs.sd	48.852	49.772	-0.094	1.226	0.027	0.087	1.023

**Sample sizes**

	Control	Treated
All	172	104
Matched	104	104
Unmatched	68	0
Discarded	0	0

**Propensity score distributions of unmatched and matched samples**

	n	mean	sd	median	trimmed	mad	min	max	range
Unmatched	68	0.11	0.04	0.11	0.11	0.05	0.03	0.17	0.14
Matched	208	0.46	0.23	0.43	0.45	0.26	0.1	0.99	0.89



Note: 0=unmatched sample; 1=matched sample

```
## glm(formula = i.aqoc ~ pg, family = binomial(link = "logit"),
##      data = dta_m)
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)   0.5521     0.2036   2.711  0.00671 **
## pg           0.7069     0.3119   2.266  0.02343 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 251.70  on 207  degrees of freedom
## Residual deviance: 246.44  on 206  degrees of freedom
## AIC: 250.44
##
## Number of Fisher Scoring iterations: 4
```