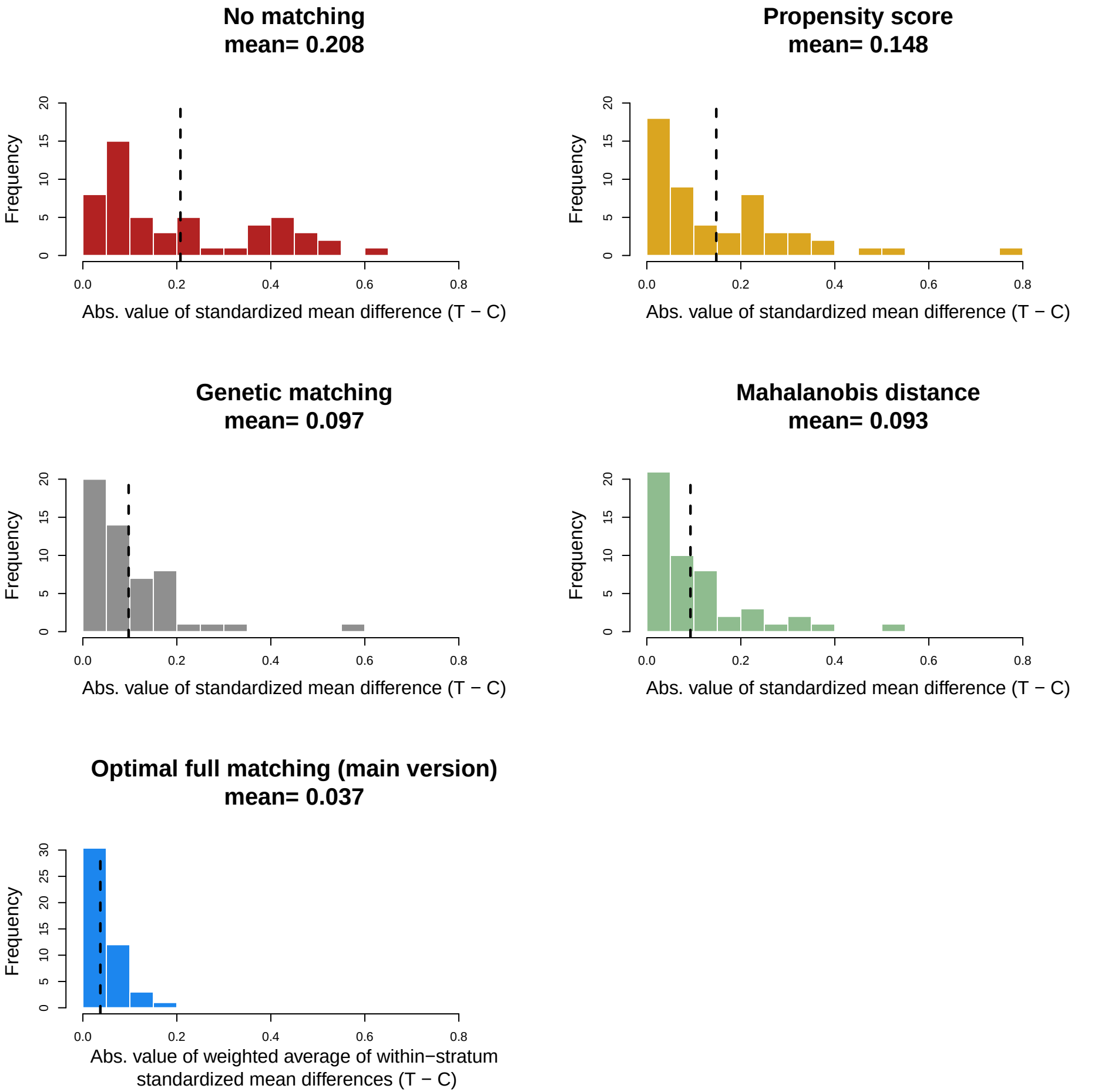
No matching
mean= 0.208
Frequency
0
5
10
15
20
0.0
0.2
0.4
0.6
0.8
Abs. value of standardized mean difference (T − C)
Propensity score
mean= 0.148
Frequency
0
5
10
15
20
0.0
0.2
0.4
0.6
0.8
Abs. value of standardized mean difference (T − C)
Genetic matching
mean= 0.097
Frequency
0
5
10
15
20
0.0
0.2
0.4
0.6
0.8
Abs. value of standardized mean difference (T − C)
Mahalanobis distance
mean= 0.093
Frequency
0
5
10
15
20
0.0
0.2
0.4
0.6
0.8
Abs. value of standardized mean difference (T − C)
Optimal full matching (main version)
mean= 0.037
Frequency
0
5
10
15
20
25
30
0.0
0.2
0.4
0.6
0.8
Abs. value of weighted average of within−stratum
standardized mean differences (T − C)