

Big__match - Testing and Demo

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This R markdown document is used for testing and demoing the current functionality of bigmatch. We'll use the sample data from the MatchIt package for basic testing.

```
library(MatchIt)
library(ggplot2)
library(ggpubr)
library(dplyr)

data("lalonde")
source('big_match.R')

# adding a binary outcome
lalonde$outcome <- lalonde$re78 > 15000
lalonde$re78 <- NULL
```

Stratify

Manual Stratify

These functions are not yet implemented.

```
# manual stratification with a continuous variable - should fail
m.strat <- manual_stratify(lalonde, "treat", "outcome",
                           covariates = c("black", "hispan", "age"))

# allowable manual stratification
m.strat <- manual_stratify(lalonde, "treat", "outcome",
                           covariates = c("black", "hispan", "nodegree"))
```

Auto Stratify

Testing Errors and Warnings

First, testing error handling. These should fail and/or give warnings.

```
# auto stratification with missing arguments  
# throws error  
a.strat <- auto_stratify(lalonde, "treat", "outcome")  
  
# auto stratification with covariates and prog scores specified, and prog_scores invalid  
# throws warning that both covariates and prog scores were specified  
# throws error that prog_scores length is invalid  
a.strat <- auto_stratify(lalonde, "treat", "outcome", c("age", "educ"), prog_scores = 1:4)
```

Testing Functionality with Valid Inputs

These should give valid results.

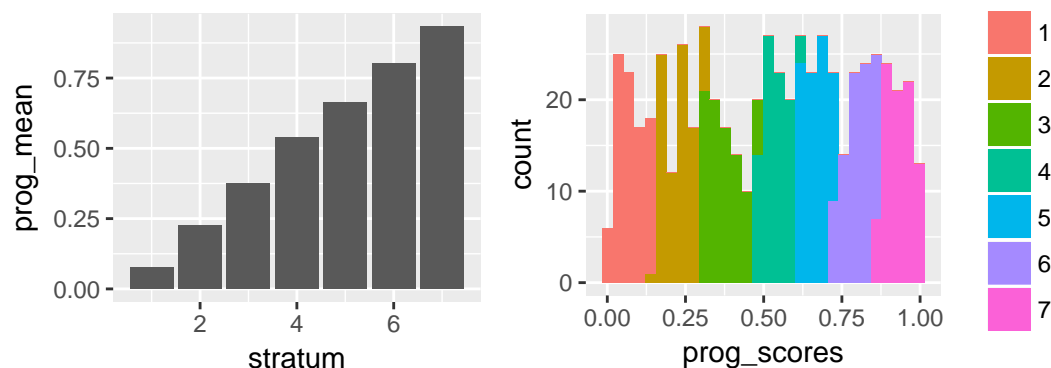
```
# auto stratification with pre-specified prognostic score  
myprogscore <- runif(n = dim(lalonde)[1])  
a.strat1 <- auto_stratify(data = lalonde, "treat", "outcome",  
                          prog_scores = myprogscore, size = 100)  
  
# auto stratification  
a.strat2 <- auto_stratify(lalonde, "treat", "outcome",  
                          covariates = c("age", "educ", "hispan", "nodegree", "black"),  
                          size = 100)  
  
# auto stratification with a non-continuous prognostic score  
a.strat3 <- auto_stratify(lalonde, "treat", "outcome",  
                          covariates = c("hispan", "nodegree", "black"), size = 100 )
```

Basic visualization of our prognostic score strata, with outcome

Below are plots of prognostic score by strata. The plot on the left shows the average prognostic score by stratum, the plot on the right gives a histogram of the prognostic scores of all samples, colored by stratum.

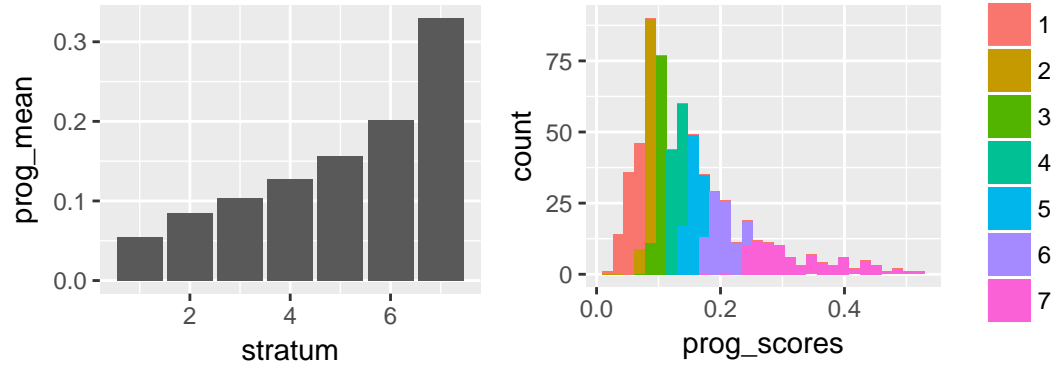
```
# uniformly generated prognostic score. Nicely continuous from 0 to 1  
basic_viz(a.strat1)
```

```
## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.
```



```
# prognostic score generated from some continuous and some discrete variables.
# Fairly continuous
basic_viz(a.strat2)
```

```
## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.
```



```
# prognostic score generated from only a few discrete variables.
# Since prog_score only takes on a few different values,
# strata quantiles are less evenly distributed from 0 to 1
basic_viz(a.strat3)
```

```
## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.
```

