

Big Match - Large Demo

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This rmarkdown is meant for running a larger demo of the `big_match` functionality.

Import Data

This sample cohort data was provided by Justin Lee at the Quantitative Sciences Unit. It contains ~900,000 observations of 112 variables.

```
dat <- read_sas("../sample_data/justincohort_june2017.sas7bdat")
```

```
# dimensions: ~900,000 x 112
dim(dat)
```

```
## [1] 893498    112
```

```
# filter and add treatment column
```

```
dat <- filter(dat, totalct > 1 & arteryCt < 3) %>%
  mutate(treat = ifelse(arteryCt > 1, 1, 0))
```

```
# dimensions: ~900,000 x 112
dim(dat)
```

```
## [1] 833657    113
```

Manual Stratify

```
m.strat <- manual_stratify(dat, treat = "treat",
                           covariates = c("Male", "race", "hosp_state"))
```

```
summary(m.strat)
```

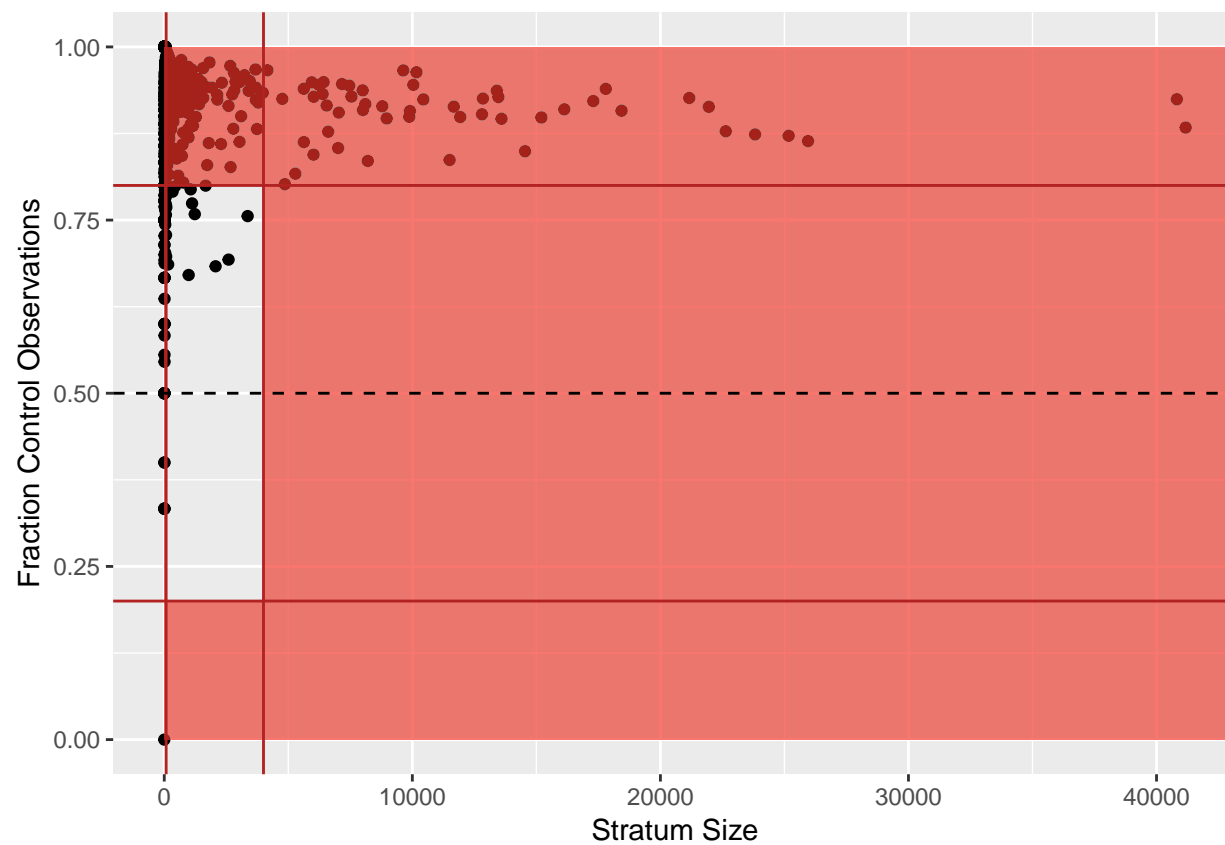
```
## $call
## manual_stratify(data = dat, treat = "treat", covariates = c("Male",
##   "race", "hosp_state"))
##
## $issue_table
## # A tibble: 704 x 6
##   Stratum Treat Control Total Control_Proporti~ Potential_Issues
##   <dbl> <dbl>   <dbl> <int>         <dbl> <chr>
## 1      1      0       3      3           1 Too few samples; Not eno~
## 2      2      0      13     13           1 Too few samples; Not eno~
## 3      3      0       6      6           1 Too few samples; Not eno~
## 4      4      1      19     20          0.95 Too few samples; Not eno~
## 5      5      0       5      5           1 Too few samples; Not eno~
## 6      6     64    1071   1135        0.944 Not enough treated sampl~
## 7      7      1      10     11          0.909 Too few samples; Not eno~
## 8      8      1      14     15          0.933 Too few samples; Not eno~
## 9      9      0       6      6           1 Too few samples; Not eno~
## 10     10      1      35     36          0.972 Too few samples; Not eno~
## # ... with 694 more rows
##
## $sum_before
##
##   Treat_Mean  Contol_Mean
##   "0"        "1"
## cohortdate   "2005-05-16" "2005-01-12"
## Admsn_source NA        NA
## adm_type     NA        NA
## BENE_ID      NA        NA
## Index_admsn_dt "2005-05-14" "2005-01-10"
## Index_dschrge_dt "2005-05-23" "2005-01-19"
## surgdate      "2005-05-16" "2005-01-12"
## PRVDR_NUM     NA        NA
## HospCABGV01_1999 NA        NA
## HospCABGV01_2000 NA        NA
## HospCABGV01_2001 NA        NA
## HospCABGV01_2002 NA        NA
## HospCABGV01_2003 NA        NA
## HospCABGV01_2004 NA        NA
## HospCABGV01_2005 NA        NA
## HospCABGV01_2006 NA        NA
## HospCABGV01_2007 NA        NA
## HospCABGV01_2008 NA        NA
## HospCABGV01_2009 NA        NA
## HospCABGV01_2010 NA        NA
## HospCABG_total  "2930.559" "2939.781"
## HospCABG_mean   "249.2153" "249.4890"
## hospname       NA        NA
```

## zipcode	NA	NA
## hosp_type	NA	NA
## urgency	"0.7812179"	"0.7172588"
## disStatus	NA	NA
## Destin	NA	NA
## hosp_state	NA	NA
## indexSurgyear	"2004.895"	"2004.558"
## deathdt	NA	NA
## CheckDeath	NA	NA
## covend	"2010-02-20"	"2010-03-18"
## dob	"1933-02-23"	"1933-12-26"
## COVSTART	"1995-11-18"	"1996-05-17"
## BENE_AGE_AT_END_REF_YR	"76.36114"	"75.60406"
## BENE_RACE_CD	NA	NA
## RTI_RACE_CD	NA	NA
## BENE_PTA_TRMNTN_CD	NA	NA
## BENE_PTB_TRMNTN_CD	NA	NA
## Male	"0.6790589"	"0.7290717"
## Sex	NA	NA
## race	NA	NA
## trmPartA	NA	NA
## TrmPartB	NA	NA
## bene_entlmt_rsn_orig	NA	NA
## bene_entlmt_rsn_curr	NA	NA
## BENE_ZIP_CD	NA	NA
## BENE_ESRD_IND	NA	NA
## BENE_MDCR_STATUS_CD	NA	NA
## BENE_HI_CVRAGE_TOT_MONS	NA	NA
## BENE_SMI_CVRAGE_TOT_MONS	NA	NA
## BENE_STATE_BUYIN_TOT_MONS	NA	NA
## BENE_HMO_CVRAGE_TOT_MONS	NA	NA
## dead	"0.2668215"	"0.2408784"
## dyear	NA	NA
## TimeTodeath_days	"1693.959"	"1847.903"
## timeTodeath_yrs	"4.637801"	"5.059297"
## ageAtdeath	NA	NA
## ageatSurgery	"72.22462"	"71.04712"
## vein_ind	" 1"	NA
## artery_ind	NA	" 1"
## RadialArtery_ind	NA	NA
## veinct	"2.469851"	"1.708752"
## arteryCt	"0.9986069"	"2.0000000"
## totalct	"3.468458"	"3.708752"
## CABGnpi1	NA	NA
## CABGmeanVol1	NA	NA
## CABGtotalVol1	NA	NA
## CABGmdname1	NA	NA
## CABGwebSpec1	NA	NA
## CABGspeccode1	NA	NA
## cabg_0thtotal1	NA	NA
## cabg_0thMean1	NA	NA
## AMI_7	"0.07465450"	"0.06983025"
## ALZH_7	"0.011032287"	"0.008417997"
## ALZH_DEMEN_7	"0.03911339"	"0.03134978"

```

## ATRIAL_FIB_7          "0.07074290" "0.06574115"
## CATARACT_7           "0.4988521"  "0.4566290"
## CHRONICKIDNEY_7      "0.1408257"  "0.1153404"
## COPD_7               "0.2069664"  "0.1758440"
## CHF_7                "0.2494214"  "0.2223765"
## DIABETES_7           "0.4151271"  "0.3851455"
## GLAUCOMA_7           "0.1479636"  "0.1378557"
## HIP_FRACTURE_7       "0.009486733" "0.007256894"
## ISCHEMICHEART_7      "0.7015238"  "0.7061652"
## DEPRESSION_7         "0.1500301"  "0.1408973"
## OSTEOPOROSIS_7      "0.06922253" "0.05506405"
## RA_OA_7              "0.3539465"  "0.3277213"
## STROKE_TIA_7         "0.1205188"  "0.1042721"
## CANCER_BREAST_7      "0.01685264" "0.01416041"
## CANCER_COLORECTAL_7 "0.02140579" "0.01851455"
## CANCER_PROSTATE_7    "0.06685648" "0.06562756"
## CANCER_LUNG_7        "0.007307581" "0.005729791"
## CANCER_ENDOMETRIAL_7 "0.002868421" "0.002498896"
## ANEMIA_7             "0.3472698"  "0.3073011"
## ASTHMA_7             "0.07162304" "0.06386067"
## HYPERL_7             "0.6875462"  "0.6787909"
## HYPERP_7             "0.1984645"  "0.1972108"
## HYPERT_7             "0.7756759"  "0.7459077"
## HYPOTH_7            "0.10364093" "0.09109611"
## Ami_Index            "0.1492971"  "0.1397993"
## cancerhistory_7      "0.1102221"  "0.1024421"
## revasc_Date          NA           NA
## Stroke_date          NA           NA
## bYPASS361_date       NA           NA
## bYPASS361_1          NA           NA
## pci_date             NA           NA
## pci_1                NA           NA
## ACUTEEmi_date        NA           NA
## ACUTEEmi_1           NA           NA
## revasc_Event         NA           NA
## stratum              "539.8588"   "563.1397"
##
## attr(,"class")
## [1] "summary.strata"
plot(m.strat)

```



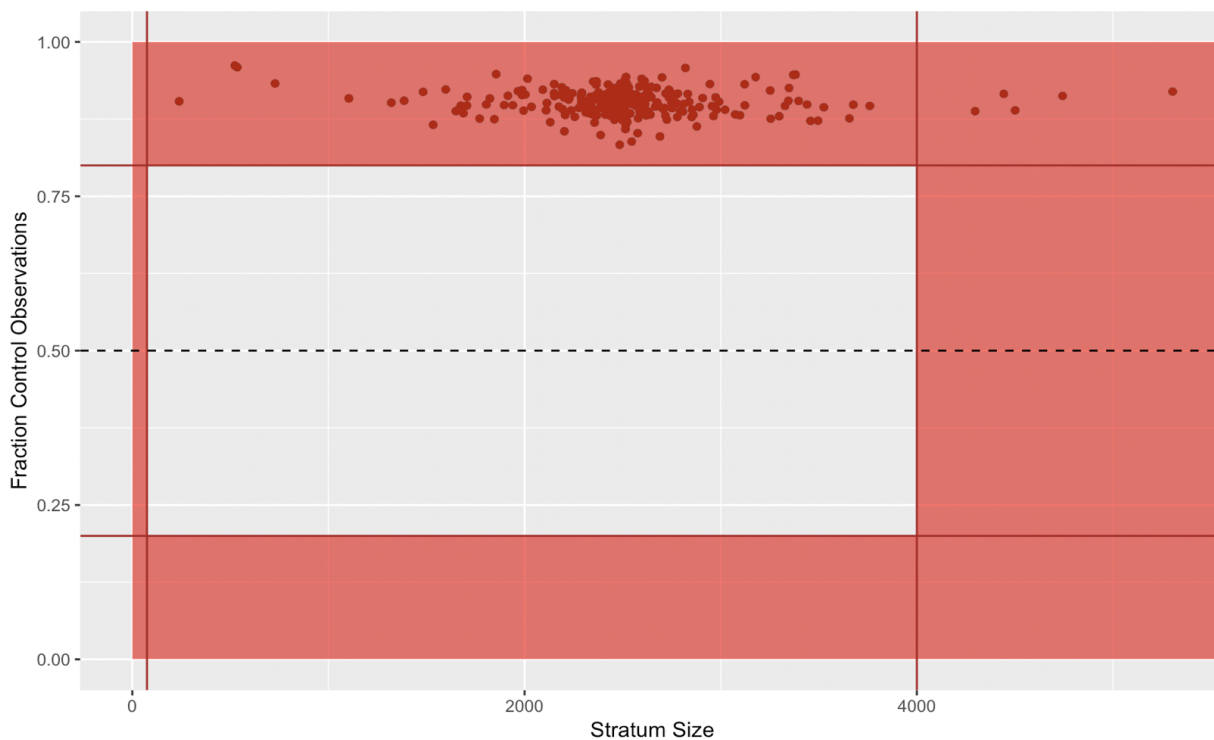


Figure 1: Auto Stratify plot.

Auto Stratify by Prognostic Score

```
a.strat <- auto_stratify(data = dat, treat = "treat",
  outcome = "dead",
  covariates = c("totalct", "hosp_state", "AMI_7", "COPD_7",
    "ISCHEMICHEART_7", "STROKE_TIA_7", "ATRIAL_FIB_7",
    "CHRONICKIDNEY_7", "DIABETES_7", "ALZH_DEMEN_7",
    "Male", "race"))

plot(a.strat)
```