

purrr::map() demo for Code review

PennCHOP microbiome program, Scott Garrett Daniel

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Packages and setting up data

Based on: <https://aosmith.rbind.io/2019/07/22/automate-model-fitting-with-loops/>

```
#devtools::install_github("thomasp85/patchwork")
library(purrr) # v. 0.3.2
library(ggplot2) # v. 3.2.0
library(patchwork) # v. 0.0.1, github only, see above ^
library(broom) # v. 0.5.2

dat = structure(list(group = structure(c(1L, 1L, 1L, 1L, 1L, 1L, 1L,
1L, 1L, 1L, 1L, 1L, 1L, 1L, 1L, 2L, 2L, 2L, 2L, 2L, 2L, 2L, 2L,
2L, 2L, 2L, 2L, 2L, 2L, 2L), .Label = c("a", "b"), class = "factor"),
  resp = c(10.48, 9.87, 11.1, 8.56, 11.15, 9.53, 8.99, 10.06,
  11.02, 10.57, 11.85, 10.11, 9.25, 11.66, 10.72, 8.34, 10.58,
  10.47, 9.46, 11.13, 8.35, 9.69, 9.82, 11.47, 9.13, 11.53,
  11.05, 11.03, 10.84, 10.22), slp = c(38.27, 46.33, 44.29,
  35.57, 34.78, 47.81, 50.45, 46.31, 47.82, 42.07, 31.75, 65.65,
  47.42, 41.51, 38.69, 47.84, 46.22, 50.66, 50.69, 44.09, 47.3,
  52.53, 53.63, 53.38, 27.34, 51.83, 56.63, 32.99, 77.5, 38.24
  ), grad = c(0.3, 0.66, 0.57, 0.23, 0.31, 0.48, 0.5, 0.49,
  2.41, 0.6, 0.27, 0.89, 2.43, 1.02, 2.17, 1.38, 0.17, 0.47,
  1.1, 3.28, 6.14, 3.8, 4.35, 0.85, 1.13, 1.11, 2.93, 1.13,
  4.52, 0.13)), class = "data.frame", row.names = c(NA, -30L) )
head(dat)
```

```
##   group  resp   slp grad
## 1     a 10.48 38.27 0.30
## 2     a  9.87 46.33 0.66
## 3     a 11.10 44.29 0.57
## 4     a  8.56 35.57 0.23
## 5     a 11.15 34.78 0.31
## 6     a  9.53 47.81 0.48
```

Function for model generation

```
ttest_fun = function(response) {
  form = paste(response, "~ group")
  lm(as.formula(form), data = dat)
}
ttest_fun(response = "resp")
```

```
##
## Call:
```

```
## lm(formula = as.formula(form), data = dat)
##
## Coefficients:
## (Intercept)          groupb
##      10.3280         -0.1207
```

Set names for vars we want to model

```
vars = names(dat)[2:4]
vars

## [1] "resp" "slp" "grad"
vars = set_names(vars)
vars

##      resp      slp      grad
## "resp"  "slp"  "grad"
```

Map the model to each var

```
models = vars %>%
  map(ttest_fun)
models

## $resp
##
## Call:
## lm(formula = as.formula(form), data = dat)
##
## Coefficients:
## (Intercept)          groupb
##      10.3280         -0.1207
##
##
## $slp
##
## Call:
## lm(formula = as.formula(form), data = dat)
##
## Coefficients:
## (Intercept)          groupb
##      43.91         4.81
```

```
##
##
## $grad
##
## Call:
## lm(formula = as.formula(form), data = dat)
##
## Coefficients:
## (Intercept)          groupb
##      0.8887          1.2773

#same thing rather than set_names in above code block
# vars %>%
#   set_names() %>%
#   map(ttest_fun)
```

Create a ggplot graph function

```
resid_plots = function(model, modelname) {
  output = augment(model) #extract residuals and fitted values via broom::augment()

  res.v.fit = ggplot(output, aes(x = .fitted, y = .resid) ) +
    geom_point() +
    theme_bw(base_size = 16)

  res.box = ggplot(output, aes(x = "", y = .resid) ) +
    geom_boxplot() +
    theme_bw(base_size = 16) +
    labs(x = NULL)

  res.v.fit + res.box +
    plot_annotation(title = paste("Residuals plots for", modelname) )
# plot_annotation is part of patchwork and glues together the individual plots
}
```

Loop through models and graph each one

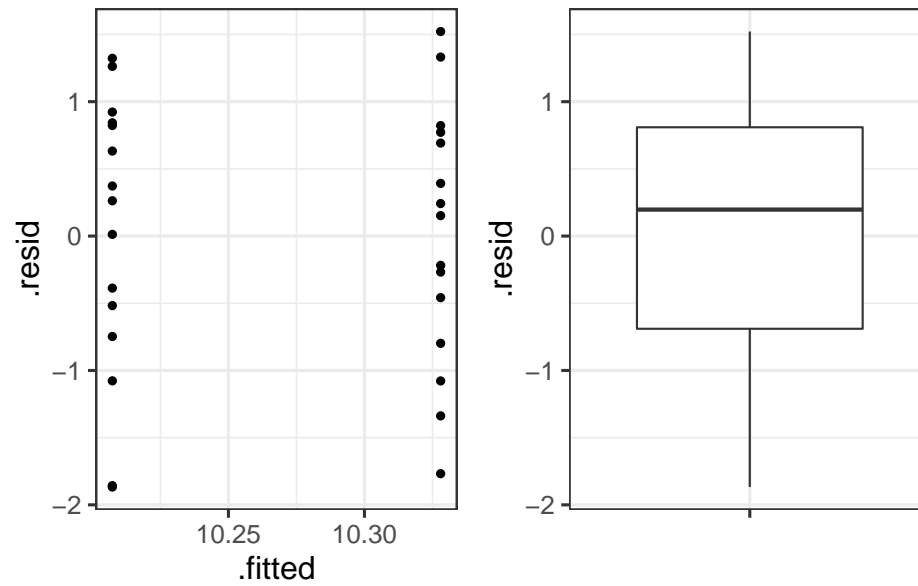
```
#for one model:
#resid_plots(model = models[[1]], modelname = names(models)[1])

#loop through all models
```

```
residplots = imap(models, resid_plots)
residplots
```

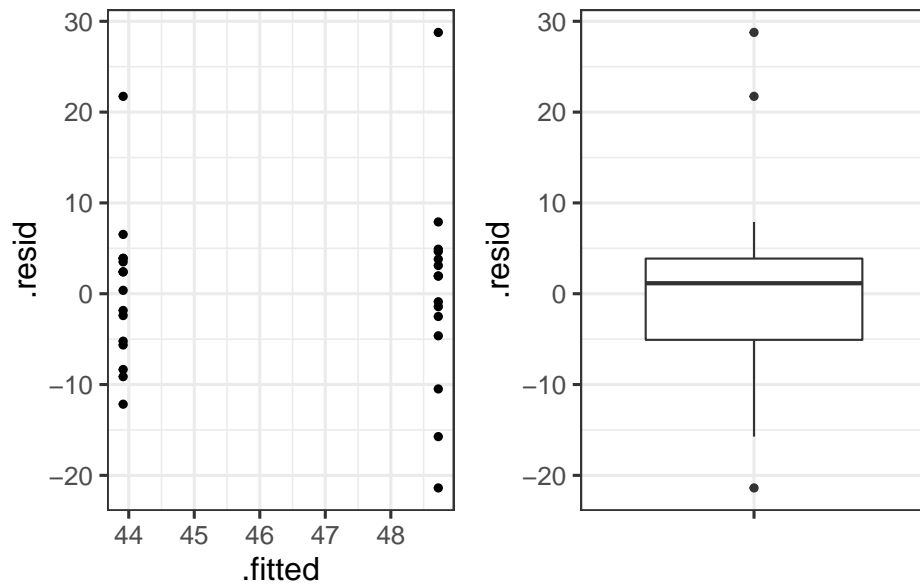
```
## $resp
```

Residuals plots for resp



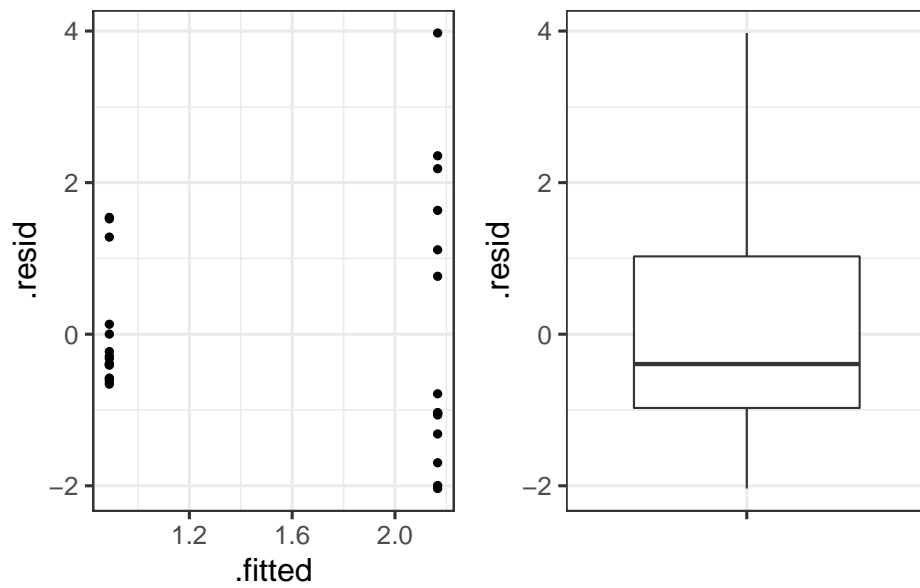
```
##  
## $slp
```

Residuals plots for slp



\$grad

Residuals plots for grad



Updating one of the models or subsetting

```
gradmod = ttest_fun("log(grad)")

models$log_grad = gradmod
models$grad = NULL
models

## $resp
##
## Call:
## lm(formula = as.formula(form), data = dat)
##
## Coefficients:
## (Intercept)      groupb
##      10.3280      -0.1207
##
##
## $slp
##
## Call:
## lm(formula = as.formula(form), data = dat)
##
## Coefficients:
## (Intercept)      groupb
##      43.91      4.81
##
##
## $log_grad
##
## Call:
## lm(formula = as.formula(form), data = dat)
##
## Coefficients:
## (Intercept)      groupb
##      -0.4225      0.7177
models[!names(models) %in% "slp"]

## $resp
##
## Call:
## lm(formula = as.formula(form), data = dat)
##
## Coefficients:
## (Intercept)      groupb
```

```
##      10.3280      -0.1207
##
##
## $log_grad
##
## Call:
## lm(formula = as.formula(form), data = dat)
##
## Coefficients:
## (Intercept)      groupb
##      -0.4225      0.7177
```

Extracting statistics from the models

```
res_anova = map_dfr(models, tidy, conf.int = TRUE, .id = "variable")

#map_dfr and map_dfc return data frames by row-binding and col-binding respectively
res_anova
```

```
## # A tibble: 6 x 8
##   variable term      estimate std.error statistic  p.value conf.low conf.high
##   <chr>    <chr>      <dbl>    <dbl>    <dbl>    <dbl>    <dbl>    <dbl>
## 1 resp    (Inter~    10.3      0.260    39.7  3.60e-26    9.80    10.9
## 2 resp    groupb    -0.121     0.368   -0.328  7.45e- 1   -0.874    0.632
## 3 slp     (Inter~    43.9      2.56    17.2  2.18e-16   38.7     49.2
## 4 slp     groupb     4.81      3.62     1.33  1.95e- 1   -2.61    12.2
## 5 log_grad (Inter~   -0.423     0.255    -1.66  1.09e- 1   -0.945    0.0997
## 6 log_grad groupb     0.718     0.361     1.99  5.64e- 2   -0.0208    1.46
```

Example from one of the projects I'm working on

Sample data

Important response and predictor variables

```
#in 16S bacterial data, we care about how the diversity changes from one condition to the next
responses = c("shannon", "richness", "faith_pd")
predictors = c("Location", "study_day")
```


Modify ttest function

```
ttest_funs = function(response, predictor) {  
  
  form = paste0(response, " ~ ", predictor)  
  lm(as.formula(form), data = s)  
  
}
```

Run it!

```
predictors = set_names(predictors,predictors)  
responses = set_names(responses,responses)  
  
for (x in predictors) {  
  responses %>%  
    map(~ ttest_funs(.,x))  
}  
  
#map2(responses, predictors, ~ ttest_funs(.x,.y))  
  
#ttest_fun(..1, ..2)
```