# Assignment 3

study group no 8

23/11/2022

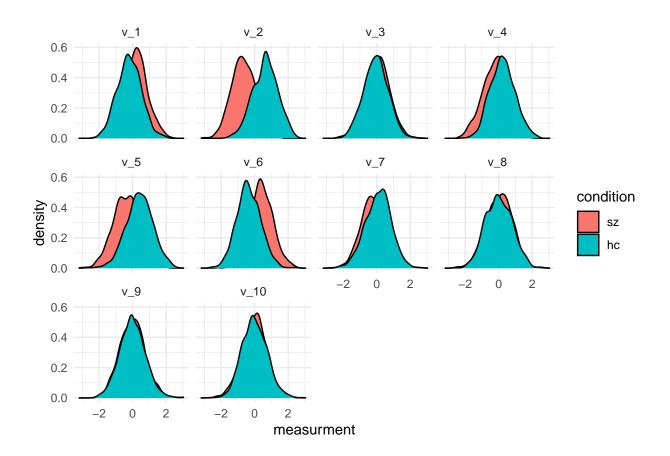
# Part I - simulating the data

Use the meta-analysis reported in Parola et al (2020), create a simulated dataset with 100 matched pairs of schizophrenia and controls, each participant producing 10 repeated measures (10 trials with their speech recorded). for each of these "recordings" (data points) produce 10 acoustic measures: 6 from the meta-analysis, 4 with just random noise. Do the same for a baseline dataset including only 10 noise variables. Tip: see the slides for the code.

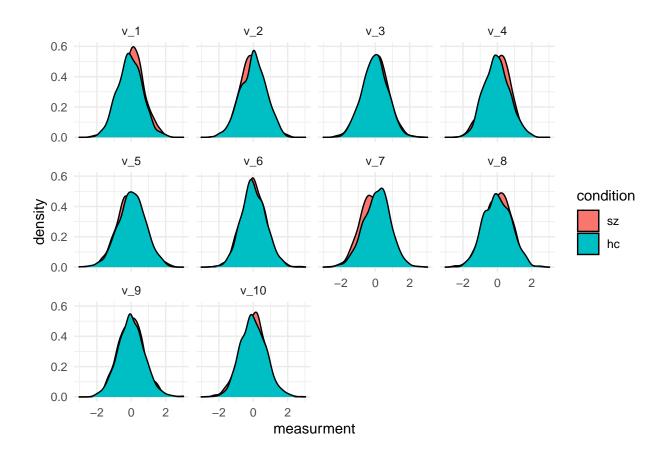
```
simulate_data <- function(pop_effects, n = 100, n_trails = 10, individual_sd = 1, trail_sd = 0.5, error
  set.seed(seed)
  tibble(
   variable = map_chr(seq_along(pop_effects), ~ paste0('v_', .)),
   population_value = pop_effects) %>%
  mutate(id = seq(1, n, by = 1) %>% list) %>%
   unnest(id) %>%
  rowwise %>%
  mutate(condition = c('sz', 'hc') %>% list,
         true_effect = rnorm(1, population_value, individual_sd) / 2,
         true_effect = c(true_effect, - true_effect) %>% list,
         trail = seq(1, n_trails, by = 1) %>% list) %>%
   unnest(c(condition, true_effect)) %>% unnest(trail) %>%
  rowwise %>%
  mutate(measurment = rnorm(1, true_effect, trail_sd) %>% rnorm(1, ., error),
         across(c(variable, id, condition), as_factor)) %>%
  relocate(c(variable, population_value), .after = condition)
m_a_values <- c(0.253, -0.546, 0.739, -1.26, -0.155, -0.75, 1.891, 0.046)
set.seed(1)
informed_pop_effects <- c(sample(m_a_values, 6, replace = F), rep(0, 4))
skeptic_pop_effects <- rep(0, 10)</pre>
dfs_long <- map(list(informed_pop_effects, skeptic_pop_effects), simulate_data)</pre>
names(dfs_long) <- c('informed', 'skeptic')</pre>
```

```
head(dfs_long[[1]])
## # A tibble: 6 x 7
## # Rowwise:
##
     id
           condition variable population_value true_effect trail measurment
##
     <fct> <fct>
                      <fct>
                                           <dbl>
                                                       <dbl> <dbl>
                                                                         <dbl>
## 1 1
           sz
                      v_1
                                           0.253
                                                      -0.187
                                                                  1
                                                                         0.603
## 2 1
                                           0.253
                                                      -0.187
                                                                  2
                                                                        -0.580
           sz
                      v_1
## 3 1
                                           0.253
                                                      -0.187
                                                                  3
                                                                        -0.485
           sz
                      v_1
## 4 1
                                           0.253
                                                      -0.187
                                                                        -0.164
           sz
                      v_1
                                                                  4
                                                                        -0.610
## 5 1
           sz
                      v_1
                                           0.253
                                                      -0.187
                                                                  5
## 6 1
           sz
                      v_1
                                           0.253
                                                      -0.187
                                                                  6
                                                                        -0.500
head(dfs_long[[2]])
## # A tibble: 6 x 7
## # Rowwise:
     id
           condition variable population_value true_effect trail measurment
##
     <fct> <fct>
                      <fct>
                                           <dbl>
                                                       <dbl> <dbl>
                                                                          <dbl>
## 1 1
                      v_1
                                               0
                                                      -0.313
                                                                         0.477
           sz
                                                                  1
## 2 1
                                               0
                                                      -0.313
                                                                        -0.706
           sz
                      v_1
                                                                  2
                                                                        -0.611
## 3 1
           sz
                      v_1
                                               0
                                                      -0.313
                                                                  3
## 4 1
                                               0
                                                      -0.313
                                                                  4
                                                                        -0.290
           sz
                      v_1
## 5 1
                                                      -0.313
           sz
                      v_1
                                               0
                                                                  5
                                                                        -0.737
## 6 1
                                                      -0.313
                                                                        -0.627
           sz
                      v_1
                                                                  6
#"'{r} #checking whether the simulation works fine
check <- map(list(informed_pop_effects, skeptic_pop_effects), ~ simulate_data(pop_effects = .x, n =
1000))
check[[1]] %>% group_by(variable) %>% summarise(mean = true_effect %>% mean, sd = true_effect
%>% sd) %>% mutate(true_mean = population_value, true_sd = 1)
check[[1]] %>% group_by(id) %>% summarise(mean = measurment %>% mean, sd = measurment %>%
sd) %>% mutate(true mean = true effect) #"'
#visualising the simulated data
map(dfs_long,
    ~ .x %>%
      ggplot(aes(x = measurment, fill = condition)) +
        geom_density()+
        facet_wrap(vars(variable)) +
        theme_minimal()
)
```

## \$informed



##
## \$skeptic



```
dfs_wide <- map(dfs_long,</pre>
 ~ .x %>%
    pivot_wider(id_cols = c(id, trail, condition),
              names_from = variable,
              values_from = measurment)
head(dfs_wide[[1]])
## # A tibble: 6 x 13
        trail condit~1 v_1 v_2 v_3
                                         v 4 v 5 v 6
                                                          v 7
    <fct> <dbl> <fct>
                     <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <
## 1 1
            1 sz
                     0.603 -1.57 -0.570 0.123 0.745 0.155 -0.156 -0.167
## 2 1
            2 sz
                     0.478
            3 sz
                     -0.485 -0.629  0.818  0.0598  0.0165  0.996  0.297
## 3 1
## 4 1
            4 sz
                     -0.164 -1.44 -0.212 0.540 0.864 0.314 0.222
                                                               0.0444
                     -0.610 -0.813 -0.516  0.425  0.283  0.237  0.0142 -1.09
            5 sz
                     6 sz
## # ... with 2 more variables: v_9 < dbl>, v_10 < dbl>, and abbreviated variable
## # name 1: condition
```

```
## # A tibble: 6 x 13
## id trail condition v_1 v_2 v_3 v_4 v_5 v_6 v_7
## <fct> <dbl> <fct> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> </dbl>
```

head(dfs wide[[2]])

```
## 1 1
                           0.477 - 0.943
                                          -0.593 0.396 1.12 -0.214 -0.156
              1 sz
## 2 1
                          -0.706 -0.913
                                          0.391 -0.374 1.02
                                                               0.582
                                                                       0.330
              2 sz
## 3 1
              3 sz
                          -0.611 0.00110 0.795 0.333 0.392 0.626
                                                                       0.297
## 4 1
              4 sz
                          -0.290 -0.806
                                         -0.235 0.813 1.24 -0.0557 0.222
## 5 1
              5 sz
                          -0.737 -0.183
                                          -0.539 0.698 0.658 -0.132
                                                                       0.0142
## 6 1
                          -0.627 -0.691
                                          0.619 -0.0336 0.710 0.212
                                                                       0.905
              6 sz
## # ... with 3 more variables: v 8 <dbl>, v 9 <dbl>, v 10 <dbl>
```

# Part II - machine learning pipeline on simulated data

On the two simulated datasets (separately) build a machine learning pipeline: i) create a data budget (e.g. balanced training and test sets); ii) pre-process the data (e.g. scaling the features); iii) fit and assess a classification algorithm on the training data (e.g. Bayesian multilevel logistic regression); iv) assess performance on the test set; v) discuss whether performance is as expected and feature importance is as expected.

Bonus question: replace the bayesian multilevel regression with a different algorithm, e.g. SVM or random forest (but really, anything you'd like to try). ## Budgeting the data:

```
# We know that using map() for a list of 2 elements might probably be considered an overkill, but we th
splits <- map(dfs_wide, ~ .x %>% initial_split(prop = 4/5))

dfs_training <- map(splits, ~ .x %>% training)
dfs_testing <- map(splits, ~ .x %>% testing)

rm(splits)
```

### Preprocessing the data

### Fitting (training) the models

### Creating the models

### Workflows

### Model fitting

### Convergance checks

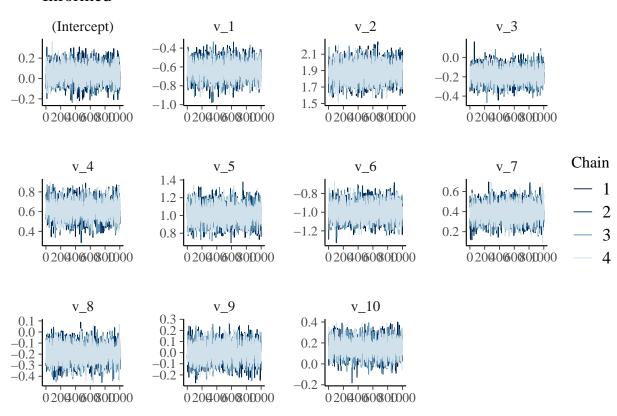
```
convergance_plots <- map2(
  fitted_models,
  names(fitted_models),
  function(.x, .y){</pre>
```

```
list(
    plot(.x, 'trace'),
    plot(.x, 'neff'),
    plot(.x, 'rhat')
    ) %>%
    map(function(.x){.x + ggtitle(.y)})
}

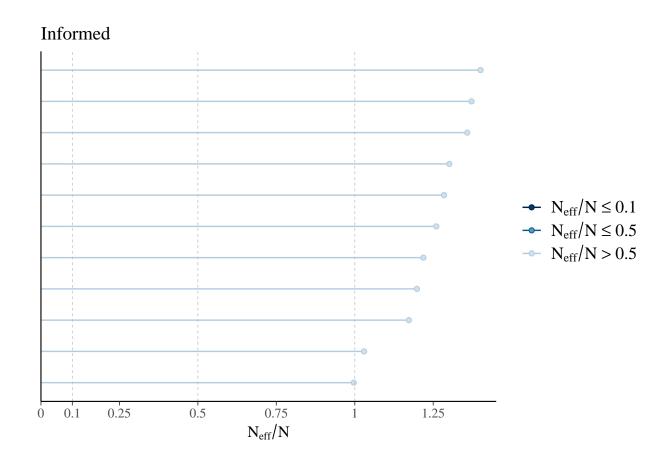
convergence_plots %>% print
```

## \$Informed
## \$Informed[[1]]

# Informed



## \$Informed[[2]]

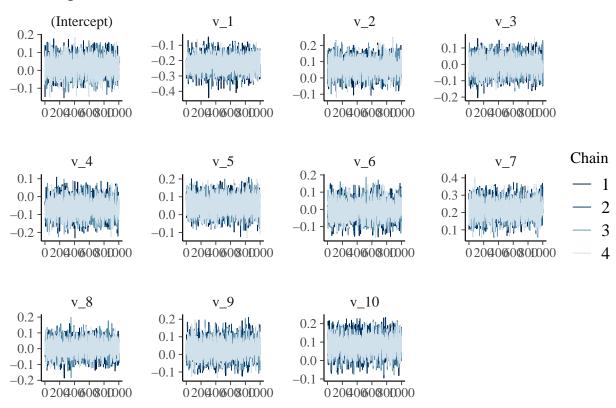


## ## \$Informed[[3]]

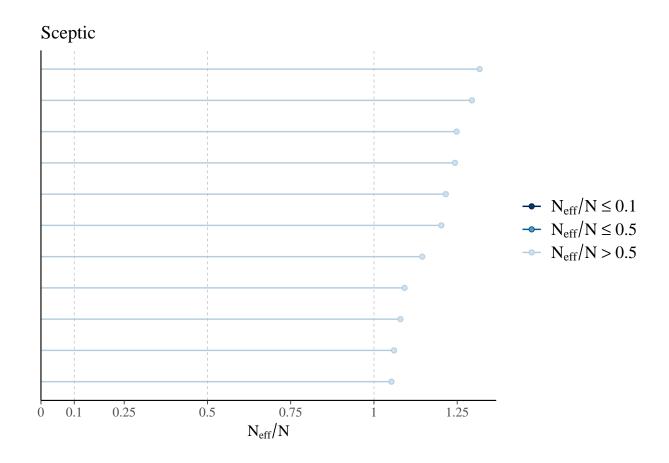
# Informed $\stackrel{\bullet}{\sim} \hat{R} \leq 1.05$ $\stackrel{\bullet}{\sim} \hat{R} \leq 1.1$ $\stackrel{\bullet}{\sim} \hat{R} > 1.1$

## ## \$Sceptic ## \$Sceptic[[1]]

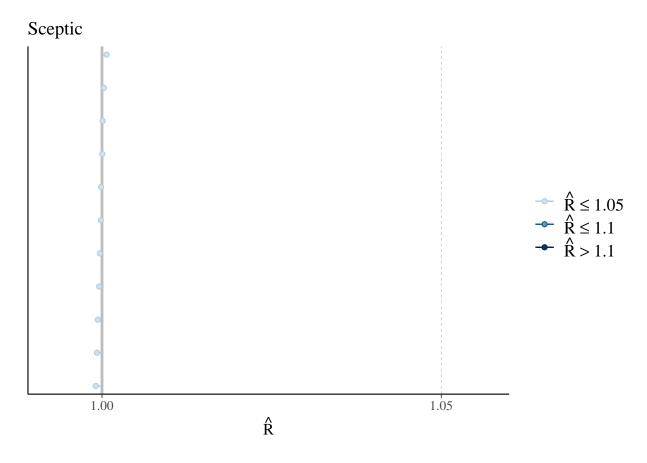
# Sceptic



## ## \$Sceptic[[2]]



## ## \$Sceptic[[3]]



```
rm(convergance_plots)
```

### Checking the priors

```
#make prior visualisation like the one on the slides (week 10)
```

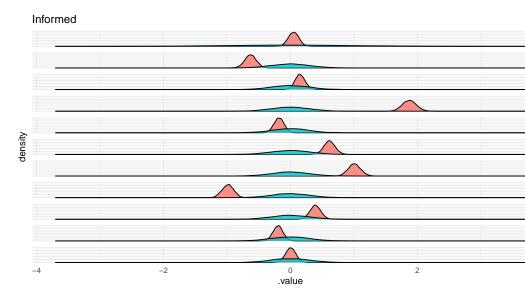
## Visualising the prior distributions

```
pp_update_plot <- function(prior_model, posterior_model){
    df_draws <-
        bind_rows(
        bind_rows(
            prior_model %>% gather_draws('(Intercept)'),
            prior_model %>% gather_draws('v_.*', regex = T)
        ) %>%
        mutate(type = 'prior'),

    bind_rows(
        posterior_model %>% gather_draws('(Intercept)'),
```

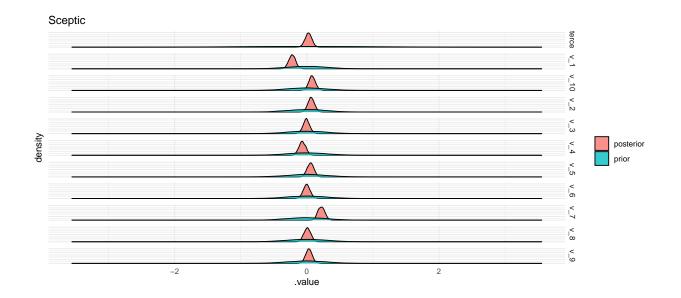
```
posterior_model %>% gather_draws('v_.*', regex = T)
        ) %>%
       mutate(type = 'posterior')
      )
  df_draws <- df_draws %>%
    group_by(.variable) %>%
    mutate(upp_lim = if_else((max(.value) + min(.value)) > 0, max(.value), - min(.value)),
           low_lim = - upp_lim) %>%
    ungroup
  df_draws %>%
    ggplot(aes(x = .value, fill = type)) +
      geom_density(alpha = 0.8) +
      labs(fill = element_blank()) +
      xlim(df_draws$low_lim[[1]], df_draws$upp_lim[[1]]) +
      facet_grid(vars(df_draws$.variable)) +
      theme_minimal() +
      theme(axis.ticks.y = element_blank(),
            axis.text.y = element_blank())
}
```

```
pp_update_plot(prior_models[[1]], fitted_models[[1]])+
   ggtitle('Informed')
```



# Prior-posterior update checks

```
pp_update_plot(prior_models[[2]], fitted_models[[2]])+
   ggtitle('Sceptic')
```



# Visualising the model

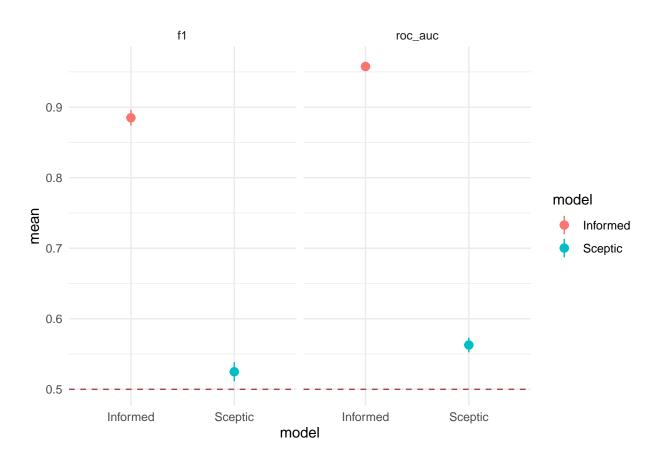
```
#add a plot of the regression line on the log-odds scale and on the probability scale
```

# Accessing model performance

### Cross-validation

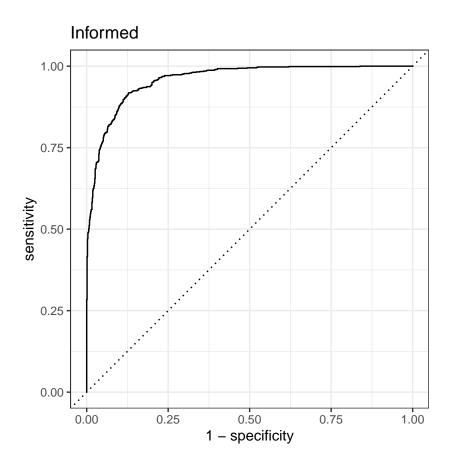
```
dfs_folded <- map(dfs_training, ~ vfold_cv(.x, v = 8))</pre>
cv_data <- map2(wflows, dfs_folded, ~ fit_resamples(.x, .y, metrics = metric_set(f_meas, roc_auc)))</pre>
cv_results <- map(cv_data, ~ collect_metrics(.x) %>%
                    mutate(upper = mean + std_err,
                           lower = mean - std_err))
cv results <- bind rows(</pre>
    cv_results[[1]] %>% mutate(model = 'Informed'),
    cv_results[[2]] %>% mutate(model = 'Sceptic')
  )
cv_results <- cv_results %>%
 rename_with(.cols = everything(), ~ str_remove(.x, stringr::fixed("."))) %>%
 mutate(metric = if_else(metric == 'f_meas', 'f1', metric))
cv_results%>%
 ggplot(aes(x = mean, y = model, xmax = upper, xmin = lower, colour = model)) +
    geom pointrange()+
    facet_wrap(vars(metric)) +
```

```
geom_vline(xintercept = 0.5, colour = 'darkred', linetype = 'dashed', alpha = 0.7) +
theme_minimal() +
coord_flip()
```

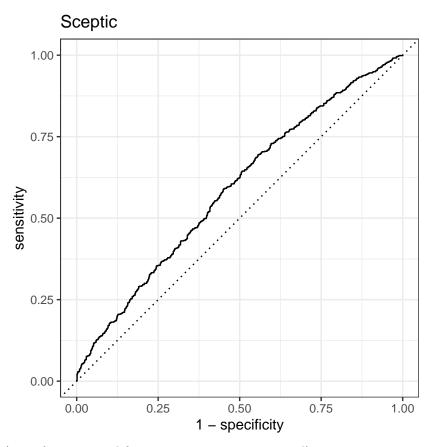


### Test data

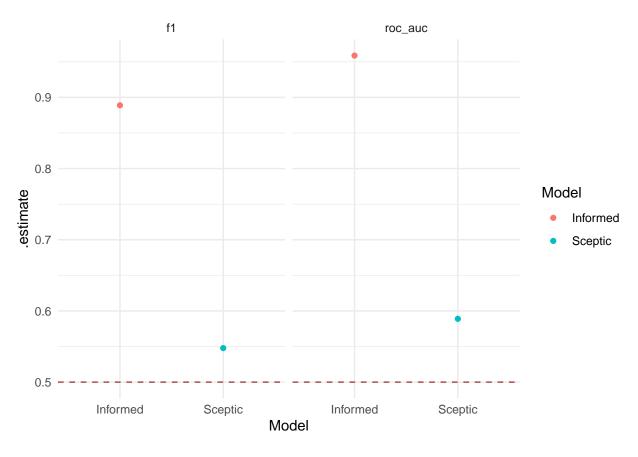
## \$Informed



## ## \$Sceptic

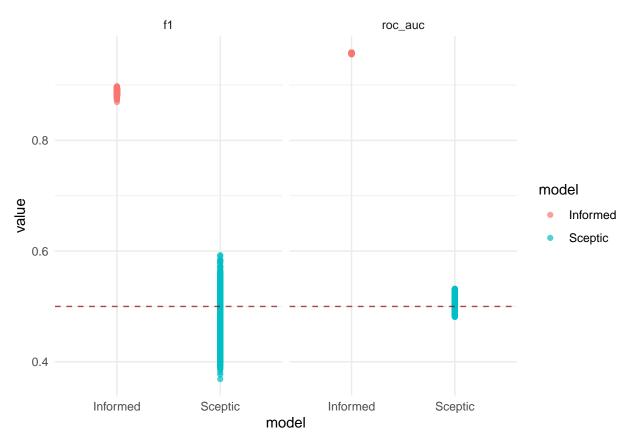


## Conclusions (is performance and feature importance as expected)



```
#with the uncertanity
test_results <- tibble(draw = NULL,</pre>
                         f1 = NULL,
                         model = NULL)
for (i in seq_along(fitted_models)){
  m <- fitted_models[[i]]</pre>
  name <- names(fitted_models)[[i]]</pre>
  draws_matrix <- posterior_epred(m)</pre>
  roc_aucs <- map_dbl(</pre>
    draws_matrix %>% split(row(draws_matrix)),
    roc_auc_vec(truth = dfs_training[[1]]$condition, estimate = .x)
  roc_aucs <- tibble(</pre>
    value = roc_aucs,
    metric = 'roc_auc',
    draw = seq_along(nrow)
    )
```

```
preds_class <- map(</pre>
    draws_matrix %>% split(row(draws_matrix)),
    ~ if_else(.x < 0.5, 'sz', 'hc') %>% as_factor %>% relevel('sz')
  fs <- map_dbl(</pre>
   preds_class,
    ~ f_meas_vec(truth = dfs_training[[1]]$condition, estimate = .x, beta = 1)
  fs <- tibble(
   value = fs,
    metric = 'f1',
    draw = seq_along(nrow)
  test_results <- bind_rows(</pre>
    test_results,
    bind_rows(fs, roc_aucs) %>% mutate(model = name)
  )
rm(i, m, name, draws_matrix, roc_aucs, preds_class, fs)
test_results <- test_results %>%
  mutate(value = if_else(metric == 'roc_auc', 1 - value, value))
test_results_summary <- test_results %>%
  group_by(model, metric) %>%
  summarise(mean = mean(value), std_err = sd(value),
            #because we're dealing the the estimates of the population parameters, the sd already is th
            lower = mean - 1.96*std_err,
            upper = mean + 1.96*std_err)
## 'summarise()' has grouped output by 'model'. You can override using the
## '.groups' argument.
test_results %>%
    ggplot(aes(x = model, y = value, colour = model)) +
      geom point(alpha = 0.7) +
      geom_hline(yintercept = 0.5, color = 'darkred', linetype = 'dashed', alpha = 0.7) +
      theme_minimal() +
      facet_wrap(vars(metric))
```

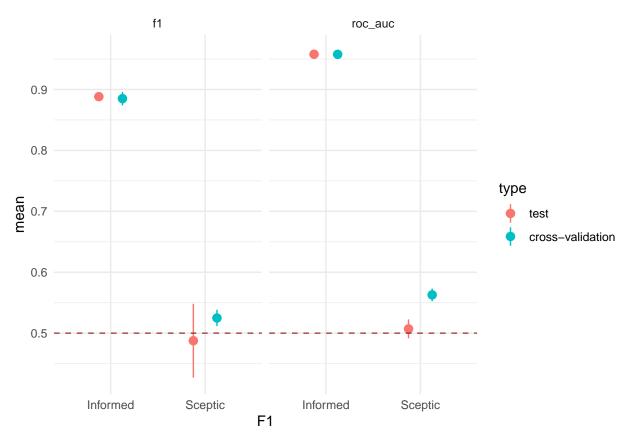


```
# Just realised this might actually not work
  # 1. mean accuracy of all draws is something very different from the accuracy of the mean linear pred
  #2. Second problem is that the confidence intervals in cross-validation and test might not show the s
        # What to do about it?
          # - plot only the accuracies only for the mean + ci of final model estimates?
          # - just back out of the confidence intervals and do all the dots for cross-validation as wel
                # - you then have to code the cross-validation 'by hand'
performance_data <- bind_rows(</pre>
 test_results_summary %>% mutate(type = 'test'),
  cv_results %>% mutate(type = 'cross-validation')) %>%
 ungroup
performance_data <- performance_data %>%
  mutate(across(where(is.character), as_factor))
glimpse(performance_data)
## Rows: 8
## Columns: 10
## $ model
               <fct> Informed, Informed, Sceptic, Sceptic, Informed, Informed, Sc~
```

<fct> f1, roc\_auc, f1, roc\_auc, f1, roc\_auc, f1, roc\_auc

## \$ metric

```
<dbl> 0.8882674, 0.9578412, 0.4875699, 0.5070287, 0.8849941, 0.957~
## $ mean
## $ std_err
               <dbl> 0.0029913341, 0.0004801837, 0.0309563668, 0.0079182479, 0.01~
               <dbl> 0.8824044, 0.9569001, 0.4268954, 0.4915089, 0.8738774, 0.952~
## $ lower
               <dbl> 0.8941305, 0.9587824, 0.5482444, 0.5225485, 0.8961109, 0.962~
## $ upper
## $ type
               <fct> test, test, test, cross-validation, cross-validation, ~
## $ estimator <fct> NA, NA, NA, NA, binary, binary, binary, binary
## $ n
               <int> NA, NA, NA, NA, 8, 8, 8
               <fct> NA, NA, NA, NA, Preprocessor1_Model1, Preprocessor1_Model1, ~
## $ config
performance_data %>%
    ggplot(aes(x = mean, y = model, xmin = lower, xmax = upper, colour = type)) +
      geom_pointrange(position = position_dodge(width = 0.5)) +
      geom_vline(aes(xintercept = 0.5), color = 'darkred', linetype = 'dashed', alpha = 0.7) +
      labs(y = 'F1') +
      theme_minimal()+
      coord_flip() +
      facet wrap(vars(metric))
```

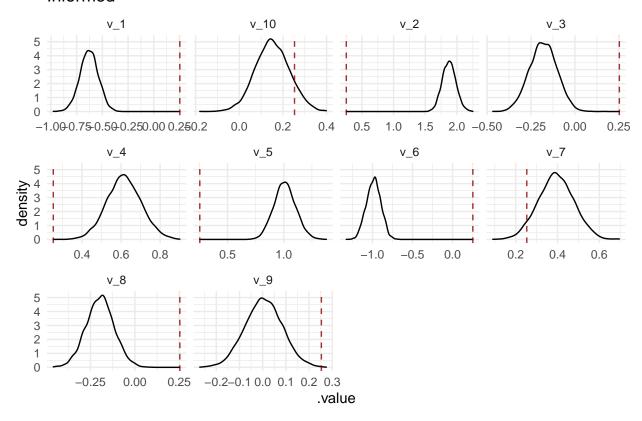


## Feature importance

```
vim_df %>%
ggplot(aes(x = .value)) +
  geom_density() +
  geom_vline(aes(xintercept = truth[[1]]), color = 'darkred', linetype = 'dashed', alpha = 0.8) +
  facet_wrap(vars(.variable), nrow = , scales = 'free_x') +
  theme_minimal()
}
vip_simulated(fitted_models[[1]], informed_pop_effects) + ggtitle('Informed')
```

## Warning: ... is ignored in group\_split(<grouped\_df>), please use group\_by(..., .add =
## TRUE) %>% group\_split()

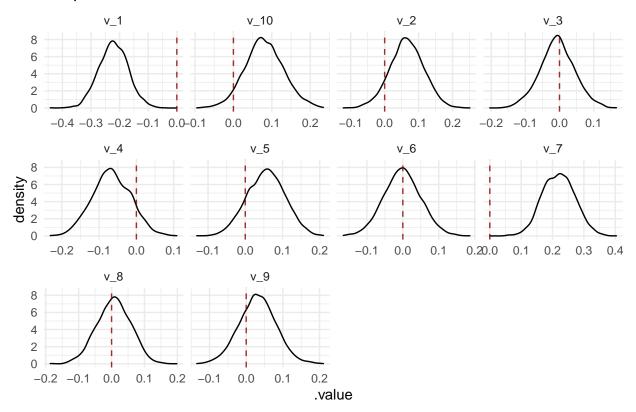
### Informed



```
vip_simulated(fitted_models[[2]], skeptic_pop_effects) + ggtitle('Skeptic')
```

## Warning: ... is ignored in group\_split(<grouped\_df>), please use group\_by(..., .add =
## TRUE) %>% group\_split()

# Skeptic

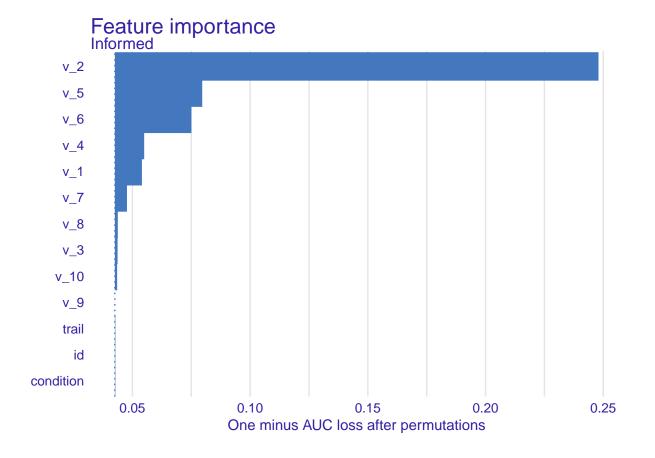


 $\#how\ to\ make\ v\_10\ appear\ as\ last?\ (mutating\ to\ factor\ before\ ggplot\ and\ inside\ facet\_wrap\ doesn't\ work)$ 

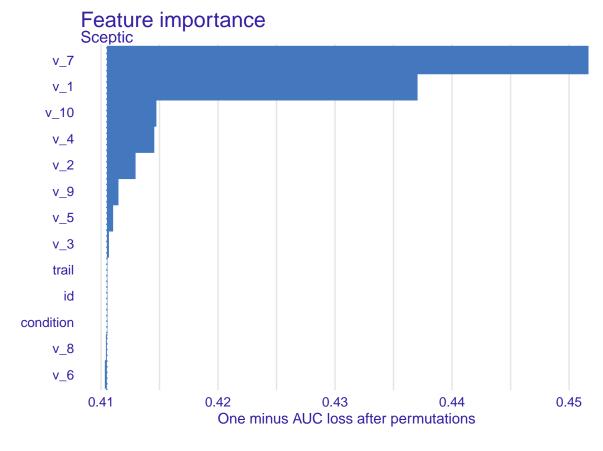
```
## Preparation of a new explainer is initiated
     -> model label
##
                         : Informed
##
     -> data
                            1600 rows 13 cols
##
     -> data
                         : tibble converted into a data.frame
                         : 1600 values
##
     -> target variable
##
     -> predict function
                        :
                            yhat.model_fit will be used ( default )
##
     -> predicted values : No value for predict function target column. ( default )
                            package parsnip , ver. 1.0.2 , task classification ( default )
##
     -> model_info
##
     -> predicted values : numerical, min = 0.000284271 , mean = 0.4993827 , max = 0.9995609
##
     -> residual function : residual_function
                         : numerical, min = 0 , mean = 0 , max = 0
##
     -> residuals
     A new explainer has been created!
## Preparation of a new explainer is initiated
```

```
-> model label : Sceptic
##
    -> data
                        : 1600 rows 13 cols
##
    -> data
                       : tibble converted into a data.frame
##
    -> target variable : 1600 values
##
    -> predict function : yhat.model_fit will be used ( default )
##
##
    -> predicted values : No value for predict function target column. ( default )
##
    -> model info
                        : package parsnip , ver. 1.0.2 , task classification ( default )
    -> predicted values : numerical, min = 0.2986658 , mean = 0.5047465 , max = 0.7008366
##
##
    -> residual function : residual_function
##
    -> residuals
                       : numerical, min = 0, mean = 0, max = 0
    A new explainer has been created!
map(
 vips,
 ~ .x %>%
   model_parts %>%
   plot(show_boxplots = F) +
     labs(title = 'Feature importance',
        subtitle = NULL)
)
```

## [[1]]

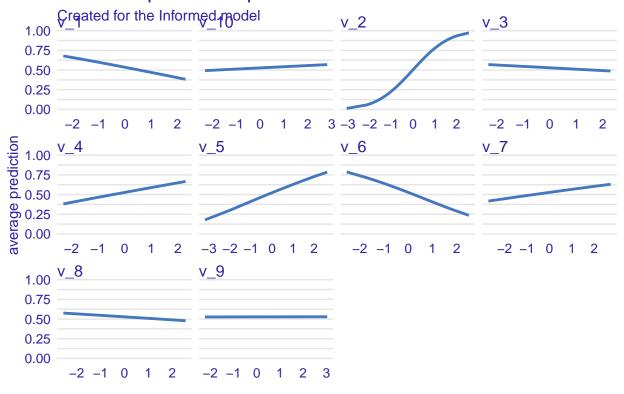


## ## [[2]]



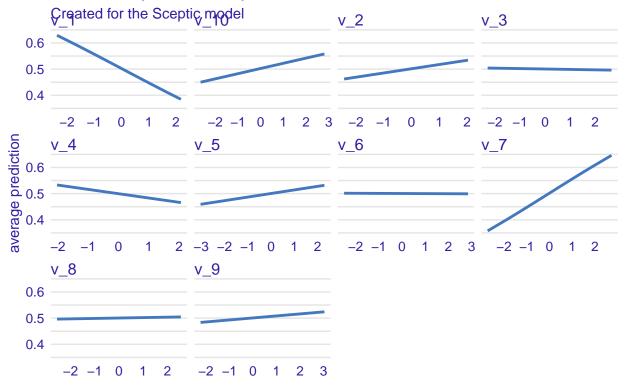
## [[1]]

# Partial dependence profile



## ## [[2]]

# Partial dependence profile



#save.image(file = "/rdata/a3\_part2.Rdata")

# Part III

Download the empirical dataset from brightspace and apply your ML pipeline to the new data, adjusting where needed. Warning: in the simulated dataset we only had 10 features, now you have many more! Such is the life of the ML practitioner. Consider the impact a higher number of features will have on your ML inference, and decide whether you need to cut down the number of features before running the pipeline (or alternatively expand the pipeline to add feature selection).

### glimpse(data\_raw)

```
## Rows: 1,889
## Columns: 398
## $ PatID
                                <chr> "101CT1", "101CT1", "101CT1", "101CT1", "~
## $ NewID
                                <chr> "CT", "CT", "CT", "CT", "CT", "CT". "CT".~
## $ Diagnosis
                                ## $ Language
                                ## $ Gender
## $ Trial
                                <chr> "T7", "T8", "T4", "T2", "T3", "T5", "T9",~
## $ Corpus
                                ## $ Duration Praat
                                <dbl> 5.62, 2.82, 9.49, 8.92, 6.00, 12.62, 10.4~
## $ FO_Mean_Praat
                                <dbl> 157.4865, 115.4691, 125.3085, 133.1547, 1~
## $ FO_SD_Praat
                                <dbl> 37.226724, 5.037427, 9.099214, 19.466738,~
## $ Intensity_Mean_Praat
                                <dbl> 70.16840, 67.47500, 70.23711, 70.42194, 6~
                                <dbl> 6.114989, 5.396695, 6.733844, 6.293483, 6~
## $ Intensity SD Praat
## $ PauseDuration_Praat
                                <dbl> 3.31, 2.00, 5.03, 5.93, 3.57, 9.22, 5.53,~
                                <dbl> 2.31, 0.82, 4.46, 2.99, 2.43, 3.40, 4.92,~
## $ TurnDuration_Praat
## $ TurnNumber_Praat
                                <dbl> 12, 5, 19, 10, 20, 23, 22, 7, 6, 9, 38, 2~
## $ PauseNumber_Praat
                                <dbl> 12, 6, 20, 10, 21, 24, 23, 8, 6, 10, 38, ~
                                <dbl> 0.4110320, 0.2907801, 0.4699684, 0.335201~
## $ PercentSpoke_Praat
## $ PercentSilence Praat
                                <dbl> 0.5889680, 0.7092199, 0.5300316, 0.664798~
## $ NHR_mean
                                <dbl> 1.0762909, 1.7420739, 0.8930114, 1.405997~
## $ NHR_std
                                <dbl> 1.0974521, 1.5076043, 1.1247343, 1.365611~
                                <dbl> 5.70, 2.90, 9.57, 9.00, 6.08, 12.70, 10.5~
## $ Duration_Cova
## $ PauseDuration_Cova
                                <dbl> 3.62, 2.10, 4.75, 5.99, 3.50, 9.26, 5.76,~
## $ TurnDuration Cova
                                <dbl> 2.08, 0.80, 4.82, 3.01, 2.58, 3.44, 4.77,~
## $ TurnNumber Cova
                                <dbl> 19, 9, 27, 28, 24, 40, 36, 15, 10, 15, 48~
                                <dbl> 19, 9, 27, 28, 24, 40, 36, 15, 10, 15, 48~
## $ PauseNumber Cova
## $ PercentSpoke Cova
                                <dbl> 0.3649123, 0.2758621, 0.5036573, 0.334444~
## $ PercentSilence_Cova
                                <dbl> 0.6350877, 0.7241379, 0.4963427, 0.665555~
## $ Pitch_Mean
                                <dbl> 5.004515, 4.823150, 4.854354, 4.878221, 4~
                                <dbl> 4.969807, 4.762174, 4.832306, 4.844187, 4~
## $ Pitch Median
## $ Pitch_SD
                                <dbl> 0.20531669, 0.17566305, 0.11439739, 0.136~
## $ Pitch_IQR
                                <dbl> 0.27026462, 0.05577301, 0.10887925, 0.115~
## $ Pitch_MAD
                                <dbl> 0.20509209, 0.03202284, 0.07263167, 0.071~
## $ F0_Mean
                                <dbl> 152.4279, 126.5938, 129.2510, 132.7043, 1~
## $ FO_Median
                                <dbl> 144.00, 117.00, 125.50, 127.00, 123.50, 1~
## $ FO SD
                                <dbl> 34.171980, 28.166559, 18.064082, 20.47695~
## $ FO_IQR
                                <dbl> 38.750, 6.625, 13.875, 15.000, 6.875, 16.~
## $ FO MAD
                                <dbl> 28.91070, 3.70650, 8.89560, 8.89560, 5.18~
## $ F1_Mean
                                <dbl> 474.3766, 590.4291, 479.5878, 539.8825, 5~
## $ F1 Median
                                <dbl> 477.8326, 596.2321, 489.8214, 504.4456, 4~
                                <dbl> 123.33146, 118.70444, 138.64037, 173.5359~
## $ F1_SD
## $ F1_IQR
                                <dbl> 156.48211, 111.39806, 150.95069, 154.2735~
                                <dbl> 119.55121, 87.83947, 109.97878, 115.86091~
## $ F1 MAD
## $ F2 Mean
                                <dbl> 1379.245, 1604.363, 1642.664, 1591.472, 1~
## $ F2_Median
                                <dbl> 1320.325, 1298.711, 1613.912, 1532.705, 1~
## $ F2 SD
                                <dbl> 359.8170, 537.6798, 442.4514, 441.9225, 4~
## $ F2 IQR
                                <dbl> 338.8416, 814.1942, 509.2859, 618.9723, 6~
## $ F2 MAD
                                <dbl> 251.7420, 284.0742, 378.3801, 450.8429, 4~
                                <dbl> 2588.769, 2725.858, 2708.497, 2575.991, 2~
## $ F3_Mean
## $ F3_Median
                                <dbl> 2668.810, 2779.451, 2682.219, 2563.081, 2~
```

```
| Adbl- | Adbl
                                                                                         <dbl> 408.1128, 398.5884, 348.2637, 348.0087, 3~
                                                                                         <dbl> 289.6221, 418.5070, 438.6633, 559.2186, 4~
                                                                                         <dbl> 171.3690, 305.8763, 341.0985, 394.4439, 3~
                                                                                        <dbl> 3378.862, 3481.532, 3584.281, 3487.865, 3~
                                                                                         <dbl> 3451.576, 3542.469, 3645.670, 3552.070, 3~
                                                                                        <dbl> 410.0392, 494.4055, 326.6898, 383.0070, 3~
                                                                                        <dbl> 590.2965, 859.3307, 412.4021, 521.6006, 3~
                                                                                         <dbl> 363.4770, 592.0502, 266.3483, 344.6552, 2~
                                                                                        <dbl> 4201.605, 4253.272, 4542.009, 4517.439, 4~
                                                                                        <dbl> 4340.185, 4365.997, 4603.801, 4651.156, 4~
                                                                                         <dbl> 456.6505, 530.5366, 325.3482, 374.0563, 2~
                                                                                         <dbl> 745.2826, 934.3287, 428.5955, 435.8510, 2~
                                                                                         <dbl> 580.2267, 546.9331, 312.7858, 242.7380, 1~
                                                                                         <dbl> 0.06180290, 0.04635425, 0.07702477, 0.043~
                                                                                         <dbl> 0.05702131, 0.03949889, 0.07097532, 0.039~
                                                                                         <dbl> 0.03384222, 0.03439789, 0.04596121, 0.030~
                                                                                         <dbl> 0.04679950, 0.04546405, 0.05426564, 0.034~
                                                                                         <dbl> 0.03489768, 0.03222384, 0.03949983, 0.025~
                                                                                        <dbl> 0.2284831, 0.1746557, 0.2585948, 0.161105~
                                                                                         <dbl> 0.2105338, 0.1607614, 0.2298884, 0.153294~
                                                                                         <dbl> 0.11634157, 0.12148572, 0.14146882, 0.088~
                                                                                        <dbl> 0.13750215, 0.12728650, 0.13913989, 0.107~
                                                                                        <dbl> 0.10112211, 0.09477261, 0.10050534, 0.077~
                                                                                        <dbl> -2.690729, -3.462465, -6.724419, -5.75338~
                                                                                       <dbl> -3.9527000, -2.8253954, -7.9101788, -6.97~
                                                                                        <dbl> 9.903949, 9.542624, 9.448744, 9.085656, 8~
                                                                                         <dbl> 14.195829, 10.139859, 12.107344, 11.26088~
                                                                                         <dbl> 10.598792, 8.434327, 9.022037, 8.413112, ~
                                                                                         <dbl> 0.4155799, 0.5096232, 0.5365308, 0.677747~
                                                                                         <dbl> 0.3856743, 0.4434876, 0.5113159, 0.654711~
                                                                                         <dbl> 0.2332902, 0.3638096, 0.2725629, 0.367708~
                                                                                         <dbl> 0.3265878, 0.6215354, 0.3690651, 0.465478~
    ## $ PSP_MAD
                                                                                        <dbl> 0.22695909, 0.40013808, 0.27969538, 0.346~
    ## $ HRF_Mean
                                                                                        <dbl> 25.63492, 26.19126, 30.71390, 31.48877, 3~
                                                                                         <dbl> 25.85260, 22.61750, 30.39679, 30.27583, 3~
    ## $ HRF_Median
    ## $ HRF SD
                                                                                        <dbl> 7.417731, 21.510680, 12.444059, 16.179379~
    ## $ HRF IQR
                                                                                        <dbl> 8.994097, 24.682586, 9.573233, 9.185110, ~
    ## $ HRF_MAD
                                                                                       <dbl> 6.761540, 18.082763, 7.055838, 7.070440, ~
                                                                                       <dbl> 0.1102170, 0.1334687, 0.1154243, 0.128729~
    ## $ MDQ Mean
                                                                                        <dbl> 0.1109119, 0.1375647, 0.1158310, 0.128661~
    ## $ MDQ_Median
   ## $ MDQ SD
                                                                                        <dbl> 0.03080837, 0.02964374, 0.03133302, 0.023~
                                                                                        <dbl> 0.04498629, 0.05090973, 0.04364999, 0.031~
   ## $ MDQ IQR
    ## $ MDQ_MAD
                                                                                        <dbl> 0.03284165, 0.03223560, 0.03258088, 0.023~
                                                    construction | c
    ## $ peakSlope_Mean
                                                                          <dbl> -0.3314857, -0.2949662, -0.3406523, -0.34~
<dbl> -0.3307617, -0.2949662, -0.3406523, -0.34~
<dbl> 0.08814333, 0.08688939, 0.09960372, 0.085~
                                                                                        <dbl> -0.3314857, -0.2978618, -0.3399992, -0.34~
   ## $ peakSlope_Median
    ## $ peakSlope_SD
   ## $ peakSlope_IQR
## $ peakSlope_MAD
    ## $ Rd_Mean
    ## $ Rd_Median
    ## $ Rd_SD
    ## $ Rd_IQR
   ## $ Rd MAD
                                                                                     <dbl> 0.5103568, 0.5689866, 0.5373712, 0.589471~
   ## $ Rd conf Mean
                                                                                        <dbl> 0.5205419, 0.5037996, 0.5403747, 0.443519~
```

```
## $ MCEP6 IQR
                       <dbl> 0.3407936, 0.3525973, 0.4895238, 0.553154~
## $ MCEP6_MAD
                       <dbl> 0.2520758, 0.2462648, 0.3665046, 0.414260~
                      <dbl> -0.4457810785, -0.0017213469, -0.40583462~
<dbl> -0.4426901477, 0.0017575745, -0.390821270~
<dbl> 0.2969627, 0.2423995, 0.3227063, 0.236961~
## $ MCEP7 Mean
## $ MCEP7 Median
## $ MCEP7 SD
## $ MCEP7_IQR
                        <dbl> 0.4169535, 0.3339511, 0.4421994, 0.300219~
```

```
## $ MCEP9_MAD

**# $ MCEP10_Mean

*** $ MCEP10_Mean

*** $ MCEP10_Sean

*** $ MCEP10_MAD

*** $ MCEP10_MAD
```

```
## $ Harmonicity Mean
                                  <dbl> 0.6462543, 0.7163654, 0.8055542, 0.709283~
## $ Harmonicity_SD
                                  <dbl> 0.4555504, 0.4600380, 0.4762750, 0.432797~
## $ Clarity Mean
                                  <dbl> 0.6575821, 0.6804585, 0.7029027, 0.671836~
                                  <dbl> 0.1593496, 0.1467127, 0.1449230, 0.131540~
## $ Clarity_SD
                                  <dbl> 6.911738, 8.040148, 7.026986, 7.208061, 6~
## $ LPerror Mean
## $ LPerror SD
                                  <dbl> 2.229282, 1.674179, 2.013750, 1.865100, 2~
## $ HarmonicProductSpectrum Mean <dbl> -13.785519, -42.346709, -9.642031, -6.450~
                                  <dbl> 92.39493, 55.07227, 94.35489, 101.06790, ~
## $ HarmonicProductSpectrum_SD
## $ CepstralPeakProminence Mean <dbl> 4.545577, 4.397983, 4.547749, 4.383870, 4~
## $ CepstralPeakProminence_SD
                                  <dbl> 0.5295720, 0.4481834, 0.6342335, 0.502741~
## $ Srh1_Mean
                                  <dbl> 0.1962962, 0.1807567, 0.1951253, 0.190560~
## $ Srh1 SD
                                  <dbl> 0.03753496, 0.03224129, 0.03797110, 0.038~
## $ Srh2_Mean
                                  <dbl> 13.321604, 5.053818, 8.798485, 13.324161,~
## $ Srh2_SD
                                  <dbl> 10.728828, 3.842785, 7.440368, 13.270542,~
## $ creakFO_Mean
                                  <dbl> 149.1352, 119.1684, 128.7470, 129.9000, 1~
## $ creakF0_SD
                                  <dbl> 28.846649, 17.551321, 15.329024, 12.77894~
## $ CreakProbability_Mean
                                  <dbl> 0.107691033, 0.322322607, 0.150356314, 0.~
## $ CreakProbability SD
                                  <dbl> 0.15440934, 0.24655873, 0.18271252, 0.169~
## $ PauseNumMin_Cova
                                  <dbl> 3.333333, 3.103448, 2.821317, 3.111111, 3~
## $ MeanPauseDur Cova
                                  <dbl> 0.19052632, 0.23333333, 0.17592593, 0.213~
## $ TurnNumMin_Cova
                                  <dbl> 3.333333, 3.103448, 2.821317, 3.111111, 3~
## $ MeanTurnDur Cova
                                  <dbl> 0.10947368, 0.08888889, 0.17851852, 0.107~
                                 <dbl> 2.135231, 2.127660, 2.107482, 1.121076, 3~
## $ PauseNumMin_Praat
## $ TurnNumMin Praat
                                 <dbl> 2.135231, 1.773050, 2.002107, 1.121076, 3~
## $ MeanTurnDur Praat
                                  <dbl> 0.1925000, 0.1640000, 0.2347368, 0.299000~
data <- data_raw %>%
 rename_with(.cols = everything(), str_to_lower) %>%
 rename(id = patid,
         condition = diagnosis) %>%
  mutate(across(where(is.character), str_to_lower),
         across(1:7, as_factor),
         condition = if_else(condition != 'ct', 'sz', 'hc') %% as_factor %% relevel('sz')) %%%
  select(-newid)
data$language %>% summary
##
      Ы
## 1889
data$corpus %>% summary
    1
        2
            3
## 681 363 375 470
data <- data %>%
  select(-language)
head(data)
```

## # A tibble: 6 x 396

```
##
           condition gender trial corpus duration~1 f0_me~2 f0_sd~3 inten~4 inten~5
     id
##
     <fct> <fct>
                    <fct> <fct> <fct>
                                                       <dbl>
                                                               <dbl>
                                                                       <dbl>
                                                                               <dbl>
                                              <dbl>
## 1 101
          hc
                     m
                            t7
                                               5.62
                                                       157.
                                                               37.2
                                                                        70.2
                                                                                6.11
## 2 101
                                                                        67.5
                            t8
                                               2.82
                                                        115.
                                                               5.04
                                                                                5.40
          hс
                     m
                                  1
## 3 101
          hc
                     m
                            t4
                                  1
                                               9.49
                                                        125.
                                                               9.10
                                                                        70.2
                                                                                6.73
## 4 101
                            t2
                                               8.92
                                                        133.
                                                              19.5
                                                                        70.4
                                                                                6.29
                                  1
          hc
                     m
## 5 101
                            t3
                                               6
                                                        122.
                                                                        67.4
                                                                                6.59
          hc
                     m
                                  1
                                                               13.5
## 6 101
                                                       132.
                                                               22.9
                                                                        69.2
                                                                                6.26
          hc
                     m
                            t5
                                  1
                                              12.6
## # ... with 386 more variables: pauseduration_praat <dbl>,
       turnduration_praat <dbl>, turnnumber_praat <dbl>, pausenumber_praat <dbl>,
       percentspoke_praat <dbl>, percentsilence_praat <dbl>, nhr_mean <dbl>,
## #
       nhr_std <dbl>, duration_cova <dbl>, pauseduration_cova <dbl>,
## #
       turnduration_cova <dbl>, turnnumber_cova <dbl>, pausenumber_cova <dbl>,
## #
       percentspoke_cova <dbl>, percentsilence_cova <dbl>, pitch_mean <dbl>,
## #
       pitch_median <dbl>, pitch_sd <dbl>, pitch_iqr <dbl>, pitch_mad <dbl>, ...
```

# Describing the data

### Condition

### Gender

```
data %>%
  count(gender) %>%
 mutate(pct = n / sum(n), pct = pct %>% round(2))
## # A tibble: 2 x 3
    gender
              n pct
    <fct> <int> <dbl>
## 1 m
            1081 0.57
## 2 f
             808 0.43
# pct should have grouped n in the denominator
data %>%
  count(gender, condition) %>%
  group_by(condition) %>%
 mutate(pct = n / sum(n), pct = pct %>% round(2))
```

```
## # A tibble: 4 x 4
## # Groups: condition [2]
    gender condition n pct
    <fct> <fct> <int> <dbl>
##
                      515 0.57
## 1 m
           SZ
## 2 m
           hc
                       566 0.57
## 3 f
                       385 0.43
           sz
## 4 f
           hc
                         423 0.43
data %>%
  count(corpus) %>%
 mutate(pct = n / sum(n), pct = pct %>% round(2))
## # A tibble: 4 x 3
##
    corpus n pct
    <fct> <int> <dbl>
            681 0.36
363 0.19
## 1 1
## 2 2
## 3 3
            375 0.2
## 4 4
             470 0.25
data %>%
  count(condition, corpus) %>%
  group_by(condition) %>%
mutate(pct = n / sum(n), pct = pct %>% round(2))
## # A tibble: 8 x 4
## # Groups: condition [2]
## condition corpus n pct
##
    <fct> <fct> <int> <dbl>
## 1 sz 1 333 0.37

## 2 sz 2 179 0.2

## 3 sz 3 151 0.17

## 4 sz 4 237 0.26

## 5 hc 1 348 0.35

## 6 hc 2 184 0.19

## 7 hc 3 224 0.23
## 8 hc
                         233 0.24
```

# Modeling the data

## Budgeting

```
data_background <- data %>% select(1:5)
data <- data %>% select(-c(gender, corpus))

split <- initial_split(data, prop = 4/5)

data_training <- training(split)
data_testing <- testing(split)

rm(split)</pre>
```

# Preprocessing the data

## Creating the models

```
prior_b <- normal(location = 0, scale = 0.3)</pre>
prior_intercept <- normal(0, 1)</pre>
model_prior <- logistic_reg() %>%
  set_engine('stan',
             prior = prior_b,
             prior_intercept = prior_intercept,
             prior_PD = T,
             cores = 3)
model <- logistic_reg() %>%
  set_engine('stan',
             prior = prior_b,
             prior_intercept = prior_intercept,
             cores = 3)
model_lasso <- logistic_reg(penalty = 0.01, mixture = 1) %>%
    set_engine('stan',
             prior = prior_b,
             prior_intercept = prior_intercept,
             cores = 3)
```

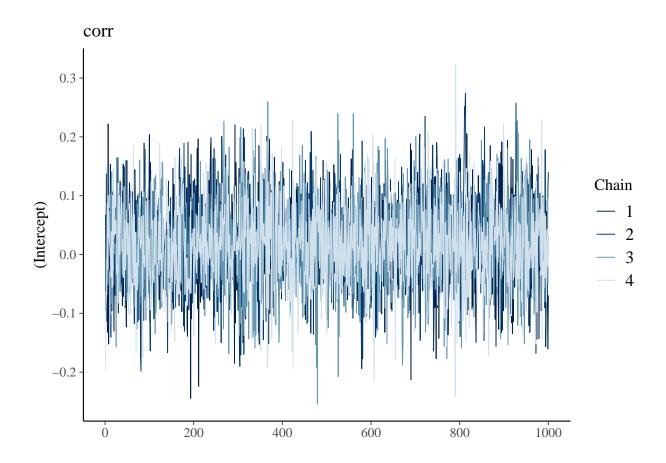
#### Workflows

## Fitting the models

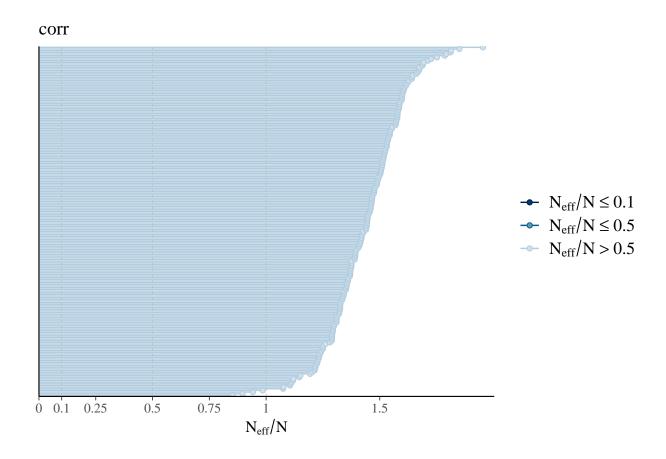
## Convergance checks

```
convergance_plots <- map2(
  fitted_models,
  names(fitted_models),
  function(.x, .y){
    list(
      plot(.x, 'trace', pars = '(Intercept)'),
      #think about which estimates to include and add this here
      plot(.x, 'neff'),
      plot(.x, 'rhat')
      ) %>%
      map(function(.x){.x + ggtitle(.y)})
  }
}
convergance_plots %>% print
```

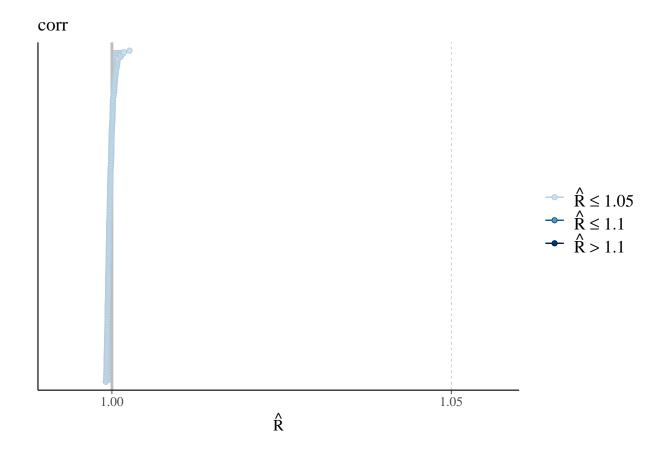
```
## $corr
## $corr[[1]]
```



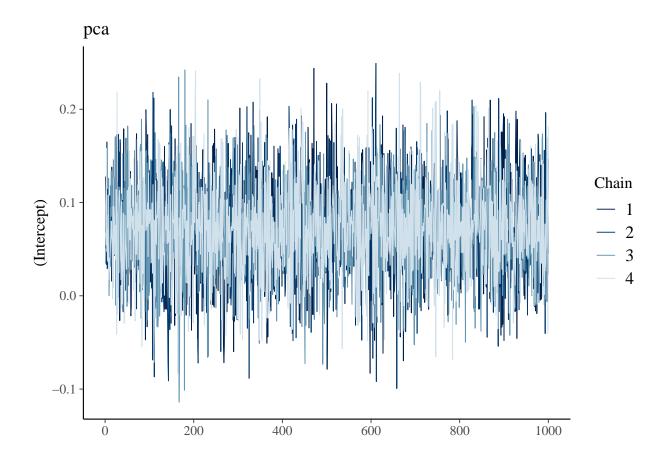
## ## \$corr[[2]]



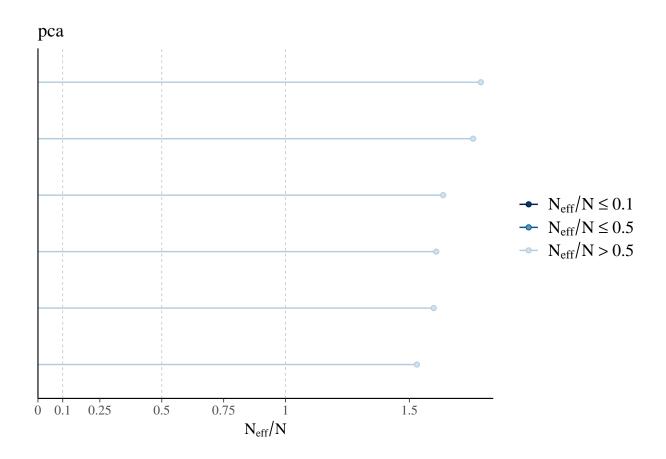
## ## \$corr[[3]]



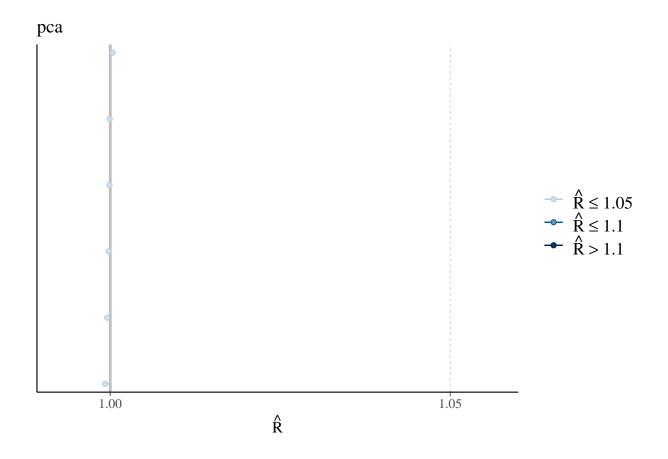
```
##
## $pca
## $pca[[1]]
```



## ## \$pca[[2]]



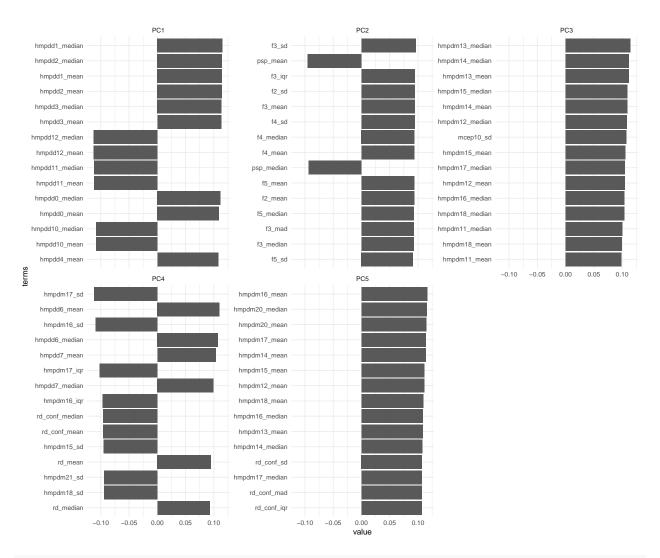
## ## \$pca[[3]]



## rm(convergance\_plots)

```
tidy_pca <- tidy(fitted[[2]] %>% extract_recipe, 2)

tidy_pca %>%
  filter(component %in% paste0('PC', 1:5)) %>%
  group_by(component) %>%
  top_n(15, abs(value)) %>%
  ungroup() %>%
  mutate(terms = reorder_within(terms, abs(value), component)) %>%
  ggplot(aes(value, terms)) +
  geom_col() +
  facet_wrap(~component, scale = 'free_y') +
  scale_y_reordered() +
  theme_minimal()
```



#qqsave('pca\_interpretation.png', height = 10, width = 12, bq = 'white')

```
variables_corr <- get_variables(fitted_models[[1]]) %>%
    str_subset('.*__', negate = T) %>%
    #removing all the 'technical' variables (e.g. 'treedepth__', 'stepsize__')
    str_subset('(Intercept)', negate = T)

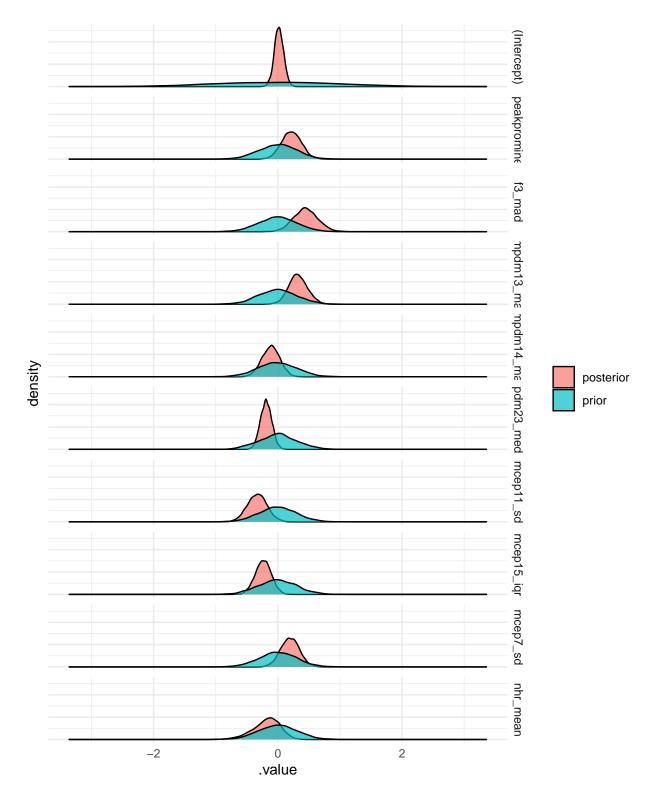
variables_corr <- c(
    '(Intercept)',
    variables_corr %>% str_subset('mcep.*') %>% sample(3, replace = F),
    variables_corr %>% str_subset('hmpdm.*') %>% sample(3, replace = F),
    variables_corr %>% str_subset(', 'mcep.*|hmpd.*', negate = T) %>% sample(3, replace = F)
)
#drawing different variables from different types of measures to plot as a sample in the pior-posterior
variables_pca <- get_variables(fitted_models[[2]]) %>% str_subset('.*__', negate = T)

pp_update_plot <- function(prior_model, posterior_model, variables){</pre>
```

df\_draws <- bind\_rows(</pre>

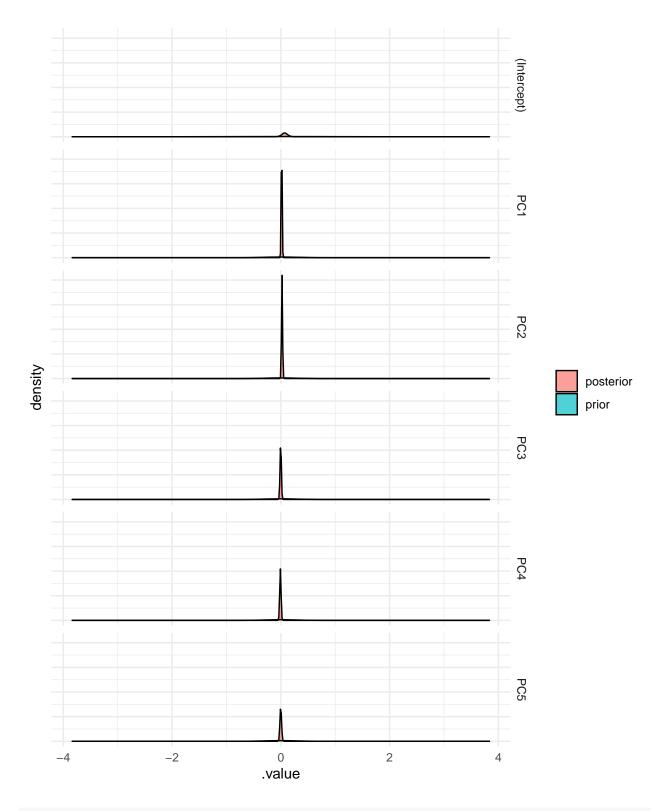
```
prior_model %>% gather_draws(!!!syms(variables))%>%
       mutate(type = 'prior'),
      posterior_model %>% gather_draws(!!!syms(variables))%>%
        mutate(type = 'posterior')
  df_draws <- df_draws %>%
    group_by(.variable) %>%
    mutate(upp_lim = if_else((max(.value) + min(.value)) > 0, max(.value), - min(.value)),
          low_lim = - upp_lim) %>%
    ungroup
  df_draws %>%
    ggplot(aes(x = .value, fill = type)) +
      geom_density(alpha = 0.7) +
      labs(fill = element_blank()) +
      xlim(df_draws$low_lim[[1]], df_draws$upp_lim[[1]]) +
      facet_grid(vars(df_draws$.variable)) +
      theme_minimal() +
      theme(axis.ticks.y = element_blank(),
            axis.text.y = element_blank())
}
```

pp\_update\_plot(prior\_fitted[[1]], fitted\_models[[1]], variables\_corr)



```
#ggsave('pp_upadte_corr.png', height = 8, bg = 'white')

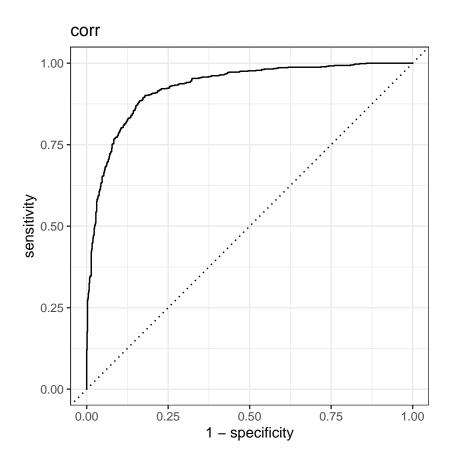
pp_update_plot(prior_fitted[[2]], fitted_models[[2]], variables_pca)
```



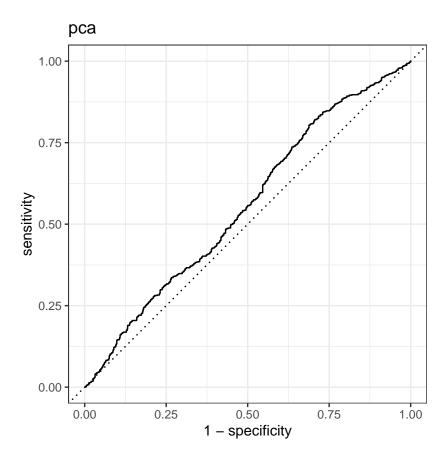
```
#ggsave('pp_update_pca.png', height = 8, bg = 'white')

test_preds <- map(fitted, ~ augment(.x, data_training))</pre>
```

## \$corr

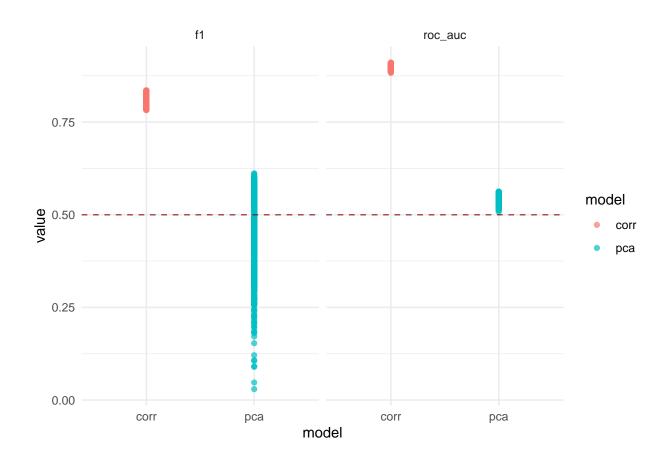


```
##
## $pca
```



```
#ggsave(plot = rocs[[1]], filename = 'roc_corr.png')
#ggsave(plot = rocs[[2]], filename = 'roc_pca.png')
```

```
preds_class <- map(</pre>
    draws_matrix %>% split(row(draws_matrix)),
    ~ if_else(.x < 0.5, 'sz', 'hc') %>% as_factor %>% relevel('sz')
 fs <- map dbl(
    preds_class,
    ~ f_meas_vec(truth = data_training$condition, estimate = .x)
  fs <- tibble(</pre>
    value = fs,
    metric = 'f1',
    draw = seq_along(nrow)
 test_results <- bind_rows(</pre>
   test_results,
    bind_rows(fs, roc_aucs) %>% mutate(model = name)
  )
}
rm(i, m, name, draws_matrix, roc_aucs, preds_class, fs)
test_results <- test_results %>%
  mutate(value = if_else(metric == 'roc_auc', 1 - value, value))
test_results %>%
    ggplot(aes(x = model, y = value, colour = model)) +
      geom_point(alpha = 0.7) +
      geom_hline(yintercept = 0.5, color = 'darkred', linetype = 'dashed', alpha = 0.7) +
      theme_minimal() +
      facet_wrap(vars(metric))
```



#ggsave('test\_results.png', bg = 'white')

 $\#save.image(file = "/rdata/a3_part3.Rdata")$