

Go to file/function

Addins

poc-recipe.R

01-dataQC.Rmd

02-preAnalytics.Rmd

03-class-POC-univariate.Rmd

04-class-POC-prelim-models.Rmd

Insert

Run

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88.1

output:
 html_document:
 code_folding: hide
 df_print: kable
 theme: flatly
 toc: yes
 toc_float: yes
params:
 my_name:
 input: text
 label: "What's your name?"
 value: "Your Name"
 reduce:
 input: select
 label: "Select which univariate output
 value: t_test
 choices: [t_test, KS, Mann_Whitney, Log
 multiple: TRUE
 stat:
 input: select
 label: "Select a statistic for feature
 value: p.value
 choices: [p.value, fdr, p.bonferroni]
 stat_cutoff:
 input: numeric
 label: "Enter a cutoff value for featur
 value: 0.1
 reduce_type:
 input: select
 label: "Do you want the intersection or
 value: intersect
 choices: [intersect, union]
 multiple: FALSE
 random_seed_repeated_cv:
 input: numeric
 label: "Enter a random seed for cross v
 value: 4321
 run_rf:
 input: checkbox

88.1 (Top Level)

Knit with Parameters

What's your name?

Your Name

Select which univariate output to use to reduce the aptamers:

t_test

Select a statistic for feature reduction:

p.value

Enter a cutoff value for feature reduction:

0.1

Do you want the intersection or the union of those selected aptamers?

intersect

Enter a random seed for cross validation:

4321

☐ Check here to run random forest model

Any CATEGORICAL clinical covariates to plot against the model residuals?

GENDER

Cancel

Knit

R Markdown