imodel_dataset

create dataset

```
linear_dgp_fun <- function(n_train, n_test, p, beta, noise_sd) {</pre>
  n <- n_train + n_test</pre>
  X <- matrix(rnorm(n * p), nrow = n, ncol = p)</pre>
  y <- X %*% beta + rnorm(n, sd = noise_sd)
  data_list <- list(</pre>
    X_train = X[1:n_train, , drop = FALSE],
    y_train = y[1:n_train],
    X_test = X[(n_train + 1):n, , drop = FALSE],
    y_{test} = y[(n_{train} + 1):n]
  return(data_list)
7
linear_dgp <- create_dgp(</pre>
  .dgp_fun = linear_dgp_fun, .name = "Linear DGP",
  # additional named parameters to pass to .dgp_fun()
  n_{train} = 350, n_{test} = 120, p = 4, beta = c(1,2,1.5,3), noise_{sd} = 1
dataset_dgp_fun <- function(datasetname){</pre>
  address <- "C:/Users/pyk/Desktop/nus/RA/project/imodels-data-master/data_cleaned/"</pre>
  file <- paste0(datasetname,".csv")</pre>
  file_path <- paste0(address,file)</pre>
  df <- read.csv(file_path)</pre>
  x \leftarrow df[, -ncol(df)]
  y <- df[, ncol(df)]
  train_indices <- createDataPartition(y, p = 0.8, list = FALSE)</pre>
```

build BART model

```
BART_fun <- function(X_train, y_train, X_test, y_test, df,k,q) {
  train_X <- data.frame(X_train)</pre>
  test_X <- data.frame(X_test)</pre>
  t <- bench::mark(fit <- wbart(x.train = train_X,
                                   y.train = y_train,
                                   x.test = test_X,
                                   k = k,
                                   sigdf = df,
                                   sigquant = q
                                   ))
  time <- mean(t$time[[1]])</pre>
  predictions <- colMeans(fit$yhat.test)</pre>
  mse_score <- mean((y_test - predictions)^2)</pre>
 return(list(time = time, mse=mse_score))
}
dbarts_fun <- function(X_train, y_train, X_test, y_test, df,k,q){</pre>
  train_X <- data.frame(X_train)</pre>
  test_X <- data.frame(X_test)</pre>
  t <- bench::mark(bart_model <- bart(x.train = train_X,
                                           y.train = y_train,
                                           x.test = test_X,
                                           k = k
                                           sigdf = df,
                                           sigquant = q))
  time <- mean(t$time[[1]])</pre>
```

```
predictions <- colMeans(bart_model$yhat.test)</pre>
     mse_score <- mean((y_test - predictions)^2)</pre>
     return(list(time = time, mse=mse_score))
}
bartMachine_fun <- function(X_train, y_train, X_test,y_test,df,k,q){</pre>
     train_X <- data.frame(X_train)</pre>
     test_X <- data.frame(X_test)</pre>
     t <- bench::mark(bart_model <- bartMachine(
                         X = train_X,
                          y = y_train,
                         k = k
                          nu = df,
                          q=q)
                     # The value of calculating the time required for modeling
     time <- mean(t$time[[1]])</pre>
     predictions <- predict(bart_model,test_X,type = "prob")</pre>
     mse_score <- mean((y_test - predictions)^2)</pre>
    return(list(time = time, mse=mse_score))
}
SoftBart_fun<- function(X_train, y_train, X_test,y_test,num_trees,alpha,beta){
     train_X <- data.frame(X_train)</pre>
     test_X <- data.frame(X_test)</pre>
     t <- bench::mark({bart_model <- softbart(X = train_X, Y = y_train, X_test = test_X, hyperations to the test_X = test_X =
                     #print(t)
     time <- mean(t$time[[1]])</pre>
     predictions <- bart_model$y_hat_test_mean</pre>
     mse_score <- mean((y_test - predictions)^2)</pre>
     return(list(time = time, mse=mse_score))
}
RF_fun <- function(X_train, y_train, X_test,y_test){</pre>
     train_X <- data.frame(X_train)</pre>
     test_X <- data.frame(X_test)</pre>
     t <- bench::mark({rf_model <- randomForest(x=train_X, y=y_train)})</pre>
     time <- mean(t$time[[1]])</pre>
     predictions <- predict(rf_model, test_X)</pre>
     mse_score <- mean((y_test - predictions)^2)</pre>
     return(list(time = time, mse=mse_score))
```

}

create evaluation

```
posterior_mse <- function(fit_results,truth_col,estimate_col){
   y_test = fit_results$truth_col
   pred = fit_results$estimate_col
   return(mean((y_test - pred)^2))
}

pred_err <- create_evaluator(
   .eval_fun = posterior_mse, .name = 'Posterior MSE',
   # additional named parameters to pass to .eval_fun()
   truth_col = "y_test", estimate_col = "predictions"
)</pre>
```

model fitting

```
BART <- create_method(</pre>
  .method_fun = BART_fun, .name = "BART",
  # additional named parameters to pass to .method_fun()
  k=2.5, q=0.95, df=4
dbarts <- create_method(.method_fun = dbarts_fun,.name = "dbarts",</pre>
                         k=2.5, q=0.95, df=4
bartMachine <- create method(.method fun = bartMachine fun,.name = "bartMachine",
                         k=2.5, q=0.95, df=4)
SoftBart <- create method(.method fun = SoftBart fun,.name = "SoftBart",</pre>
                         num_trees=50,alpha=0.95,beta=2)
RF <- create_method(.method_fun = RF_fun,.name = "RandomForest")
# Create experiment
experiment <- create_experiment(name = "Test Experiment") %>%
  add_dgp(linear_dgp) %>%
  add_dgp(dataset_dgp) %>%
  add_method(dbarts) %>%
  add method(BART) %>%
  add method(bartMachine) %>%
  add_method(SoftBart) %>%
```

```
add_method(RF)%>%
add_evaluator(pred_err)

results <- run_experiment(experiment, n_reps = 4, save = TRUE)</pre>
```

Fitting Test Experiment...

Warning: Some expressions had a GC in every iteration; so filtering is disabled.

Warning: Some expressions had a GC in every iteration; so filtering is

Warning: Some expressions had a GC in every iteration; so filtering is disabled.

```
Warning in randomForest.default(x = train_X, y = y_train): The response has
five or fewer unique values. Are you sure you want to do regression?
Warning in randomForest.default(x = train_X, y = y_train): The response has
five or fewer unique values. Are you sure you want to do regression?
Warning in randomForest.default(x = train_X, y = y_train): The response has
five or fewer unique values. Are you sure you want to do regression?
Warning in randomForest.default(x = train_X, y = y_train): The response has
five or fewer unique values. Are you sure you want to do regression?
Warning in randomForest.default(x = train_X, y = y_train): The response has
five or fewer unique values. Are you sure you want to do regression?
Warning in randomForest.default(x = train_X, y = y_train): The response has
five or fewer unique values. Are you sure you want to do regression?
Warning in randomForest.default(x = train_X, y = y_train): The response has
five or fewer unique values. Are you sure you want to do regression?
Warning in randomForest.default(x = train_X, y = y_train): The response has
five or fewer unique values. Are you sure you want to do regression?
Warning in randomForest.default(x = train_X, y = y_train): The response has
five or fewer unique values. Are you sure you want to do regression?
Warning in randomForest.default(x = train_X, y = y_train): The response has
five or fewer unique values. Are you sure you want to do regression?
Warning in randomForest.default(x = train_X, y = y_train): The response has
five or fewer unique values. Are you sure you want to do regression?
Warning in randomForest.default(x = train_X, y = y_train): The response has
five or fewer unique values. Are you sure you want to do regression?
Warning in randomForest.default(x = train_X, y = y_train): The response has
```

```
five or fewer unique values. Are you sure you want to do regression?
Warning in randomForest.default(x = train_X, y = y_train): The response has
five or fewer unique values. Are you sure you want to do regression?
Warning in randomForest.default(x = train_X, y = y_train): The response has
five or fewer unique values. Are you sure you want to do regression?
Warning in randomForest.default(x = train_X, y = y_train): The response has
five or fewer unique values. Are you sure you want to do regression?
Warning in randomForest.default(x = train_X, y = y_train): The response has
five or fewer unique values. Are you sure you want to do regression?
Warning in randomForest.default(x = train_X, y = y_train): The response has
five or fewer unique values. Are you sure you want to do regression?
Warning in randomForest.default(x = train_X, y = y_train): The response has
five or fewer unique values. Are you sure you want to do regression?
Warning in randomForest.default(x = train_X, y = y_train): The response has
five or fewer unique values. Are you sure you want to do regression?
Saving fit results...
Fit results saved | time taken: 0.053212 seconds
4 reps completed (totals: 4/4) | time taken: 4.716040 minutes
Evaluating Test Experiment...
Warning: Unknown or uninitialised column: `truth_col`.
Warning: Unknown or uninitialised column: `estimate_col`.
Evaluation completed | time taken: 0.000149 minutes
Saving eval results...
Eval results saved | time taken: 0.042850 seconds
_____
No visualizers to visualize. Skipping visualization.
_____
# Render automated documentation and view results
#render docs(experiment)
```

result <- results\$fit_results result

```
# A tibble: 40 x 5
        .dgp_name .method_name time
                                                mse
  <chr> <chr>
                   <chr>
                                t>
                                               <dbl>
1 1
        Linear DGP BART
                                <bench_tm [1]> 1.40
2 1
        Linear DGP RandomForest <bench_tm [1]> 3.53
3 1
        Linear DGP SoftBart
                              <bench_tm [1]> 1.41
4 1
        Linear DGP bartMachine <bench_tm [1]> 1.46
        Linear DGP dbarts
                              <bench_tm [1]> 1.33
6 1
        heart
                   BART
                               <bench_tm [1]> 0.130
7 1
        heart
                   RandomForest <bench_tm [1] > 0.131
8 1
                                <bench_tm [1]> 0.133
        heart
                   SoftBart
9 1
                   bartMachine <bench_tm [1]> 0.130
        heart
10 1
                                <bench_tm [1]> 1.26
        heart
                   dbarts
# i 30 more rows
```

result\$time_numeric <- as.numeric(result\$time)</pre>

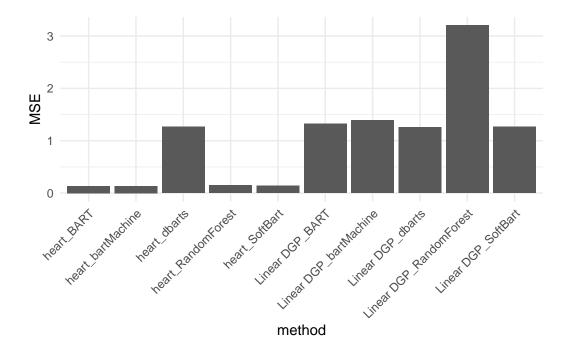
```
result$Resource <- paste(result$.dgp_name, result$.method_name, sep="_")

# Calculate MSE for each group
summary <- result %>%
    group_by(Resource) %>%
    summarise(
        Mean_MSE = mean(mse),
        Var_MSE = sd(mse),
        Mean_time = mean(time_numeric),
        Var_time = sd(time_numeric))
```

A tibble: 10 x 5

Resource	Mean_MSE	Var_MSE	${\tt Mean_time}$	Var_time
<chr></chr>	<dbl></dbl>	<dbl></dbl>	<dbl></dbl>	<dbl></dbl>
1 Linear DGP_BART	1.33	0.242	2.46	0.0338
2 Linear DGP_RandomFo	rest 3.21	0.589	0.132	0.0326
3 Linear DGP_SoftBart	1.27	0.302	15.0	0.0884
4 Linear DGP_bartMach	ine 1.39	0.207	0.792	0.0290
5 Linear DGP_dbarts	1.25	0.152	0.596	0.0160
6 heart_BART	0.135	0.00371	2.22	0.0129
7 heart_RandomForest	0.149	0.0151	0.132	0.00273
8 heart_SoftBart	0.137	0.00366	9.75	0.0517
9 heart_bartMachine	0.135	0.00355	0.617	0.0293

```
ggplot(summary, aes(x = Resource, y = Mean_MSE)) +
  geom_bar(stat = "identity", position = "dodge") +
  theme_minimal() +
  theme(axis.text.x = element_text(angle = 45, hjust = 1))+
  labs(y = "MSE", x = "method")
```



```
ggplot(summary, aes(x = Resource, y = Mean_time)) +
  geom_bar(stat = "identity", position = "dodge") +
  theme_minimal() +
  theme(axis.text.x = element_text(angle = 45, hjust = 1))+
  labs(y = "time", x = "method")
```

