

Navigating Insurance Claim Data Through Tidymodels Universe

Future of Insurance

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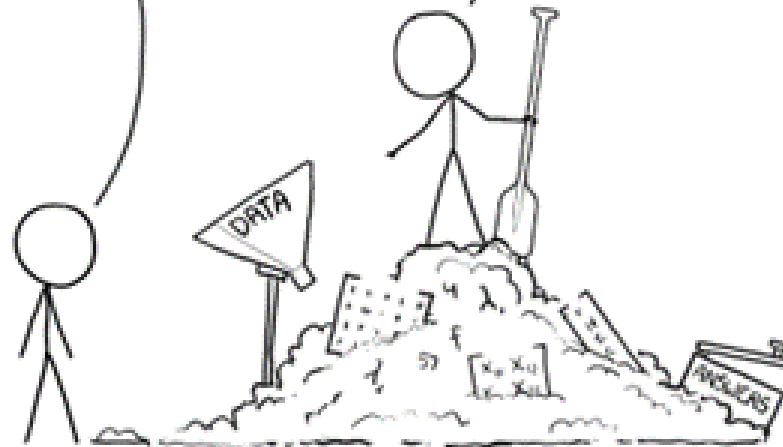
26 Aug 2021

THIS IS YOUR MACHINE LEARNING SYSTEM?

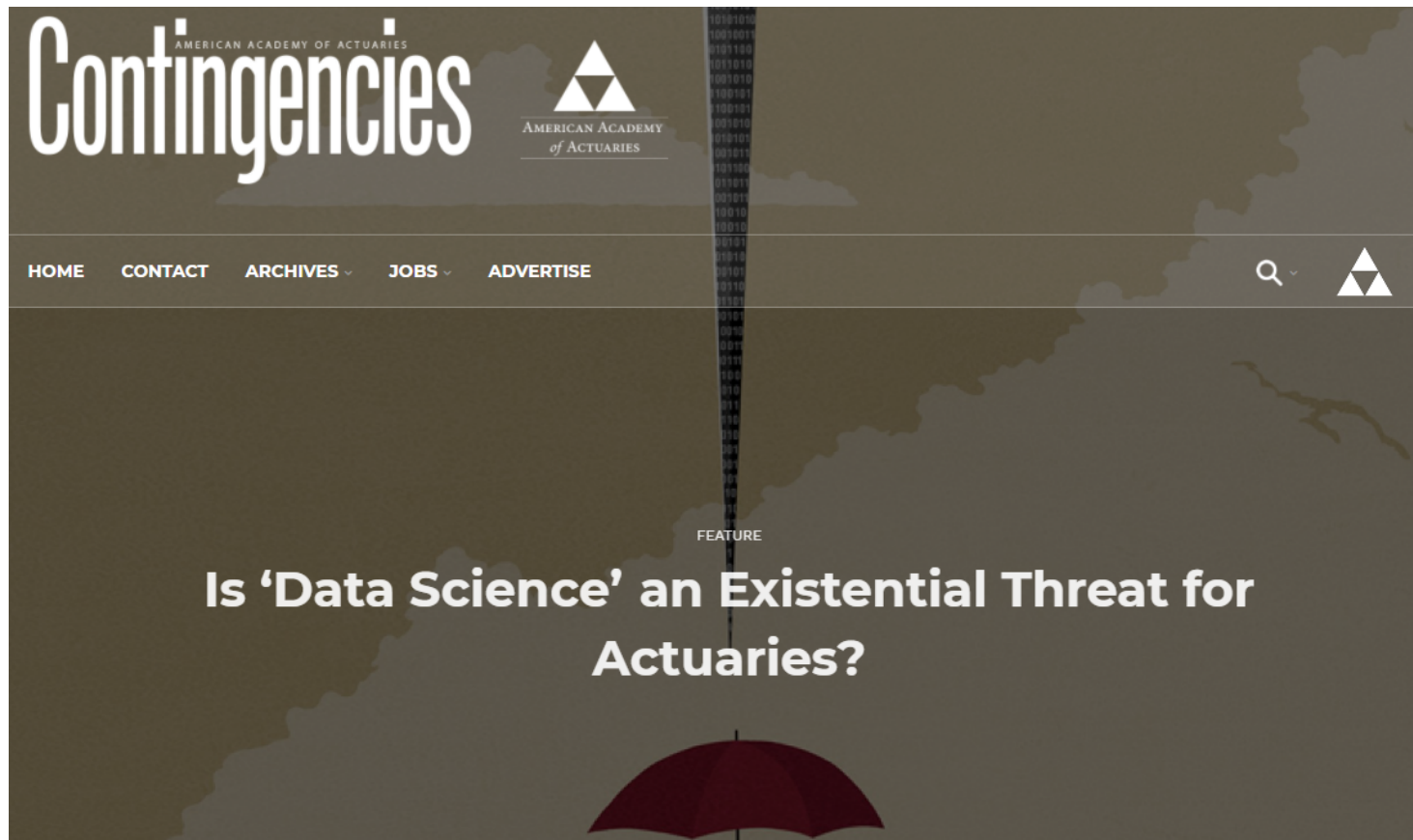
YUP! YOU POUR THE DATA INTO THIS BIG
PILE OF LINEAR ALGEBRA, THEN COLLECT
THE ANSWERS ON THE OTHER SIDE.

WHAT IF THE ANSWERS ARE WRONG?

JUST STIR THE PILE UNTIL
THEY START LOOKING RIGHT.



Are data scientists taking over actuaries?



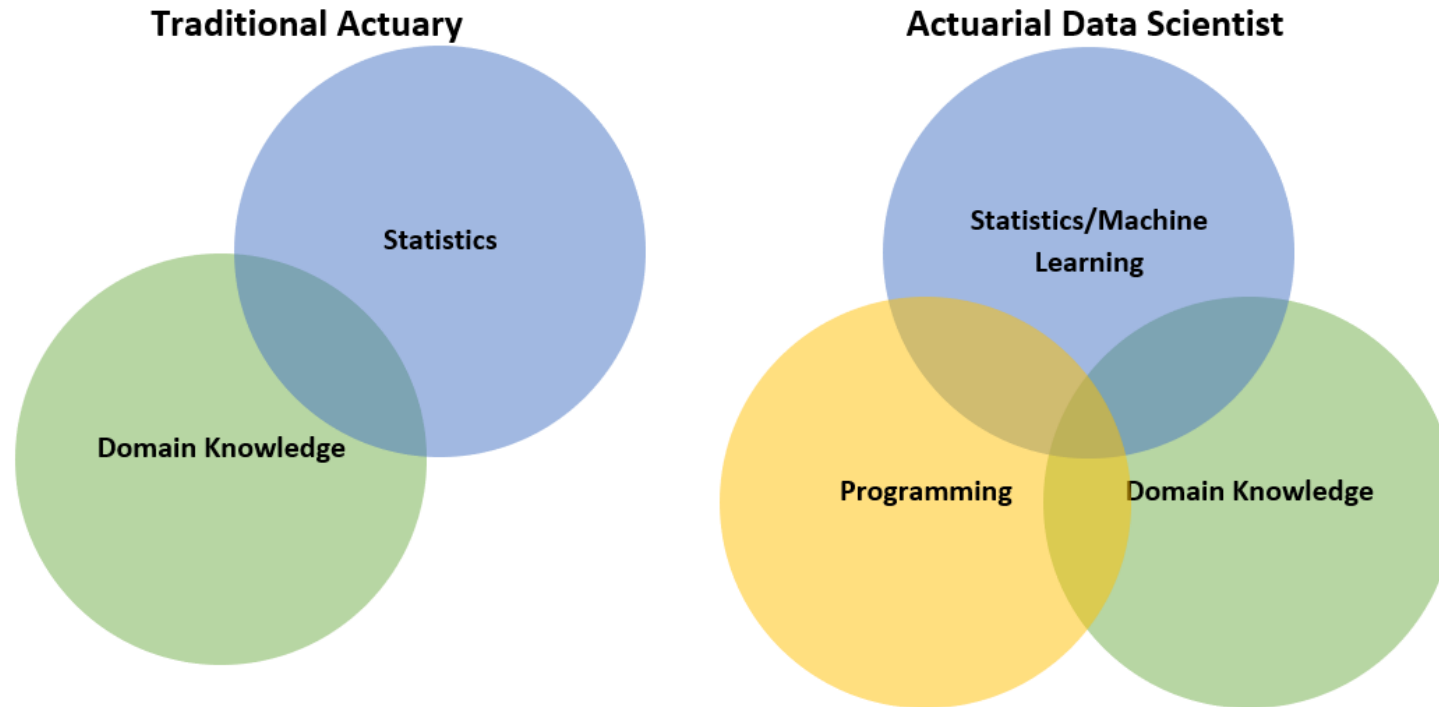
Source: *Contingencies Volume 32 No 1*

Should actuaries start looking for other jobs in other industries?

Relax! Most articles agree that data scientists are not likely to take over actuaries due to the complexity of actuarial science.

In general, these articles also pointed out there are much more actuaries could learn from data scientists to assist us in our actuarial analysis.

Additional Skillset for Actuaries to acquire



*Modified the graph by **Quantee***

It is not just about learning the programming language, but also what are the best practice of performing machine learning tasks (eg. what is the suitable package to use to perform the necessary analysis).

When 1 + 1 > 2

How Modern Data Science Could Complement Actuarial Science in Claim Cost Estimation

Jasper LOK Jun Haur Supervisor: Prof. KAM Tin Seong

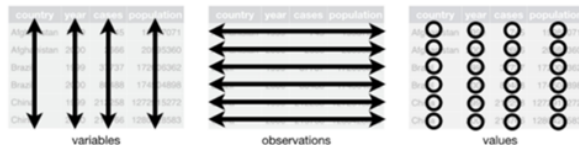
INTRODUCTION

The increasing ability to store and analyze the data due to the advancement in technology has provided actuaries opportunities in optimizing capital held by insurance companies. The modern data science packages also allows the actuaries harness the power of data science to mine the “gold” in unstructured data. This would help the companies in gaining competitive advantage in the market.

MOTIVATION

In this capstone project, various modern data science techniques are used to demonstrate how they provides the actuaries another toolkit to sharpen the conventional actuarial analysis. **tidyverse** and **tidymodels** packages are being selected for the demonstration.

TIDY DATA CONCEPT



Following are the definition of tidy data:

- Each variable must have its column
- Each observation must have its row
- Each value must have its cell

Traditional Actuary



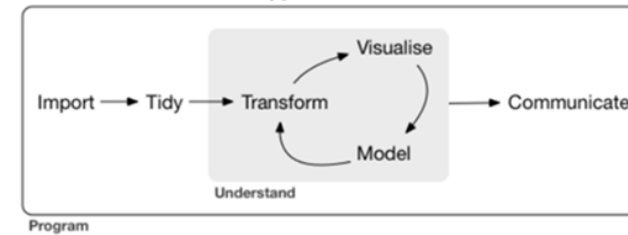
Actuarial Data Scientist



Analysis tools used:

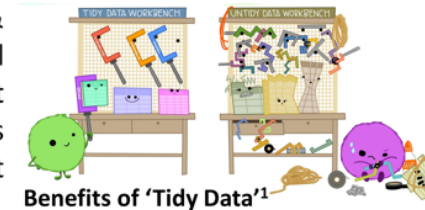


Typical Data Science Workflow:



EXPLANATORY DATA ANALYSIS (EDA)

Various R packages from **tidyverse** is used to import & clean data, check data quality and perform statistical test. The main benefit of using these packages is that they conform to tidy data concept, allowing the users effectively perform necessary data analysis, without spending too much time on data transformation.

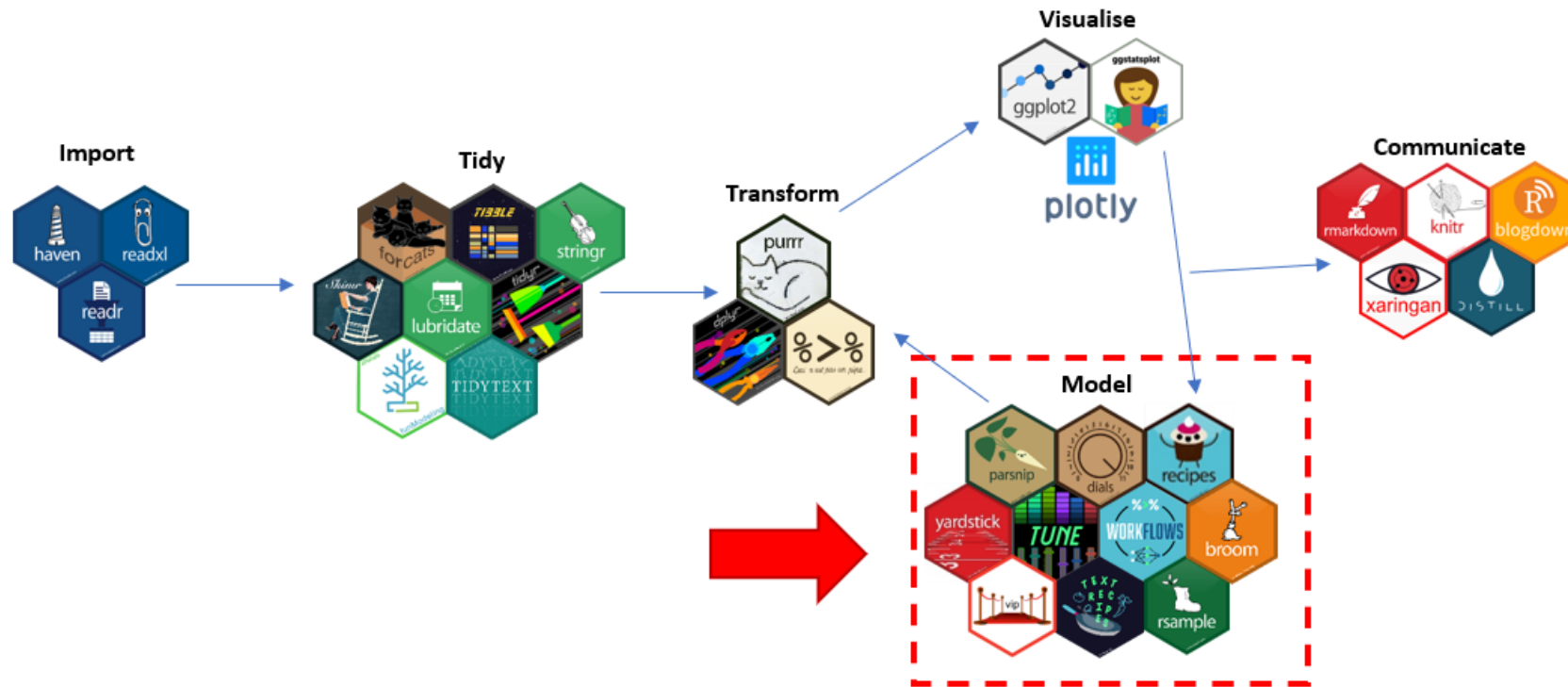


Benefits of 'Tidy Data'¹



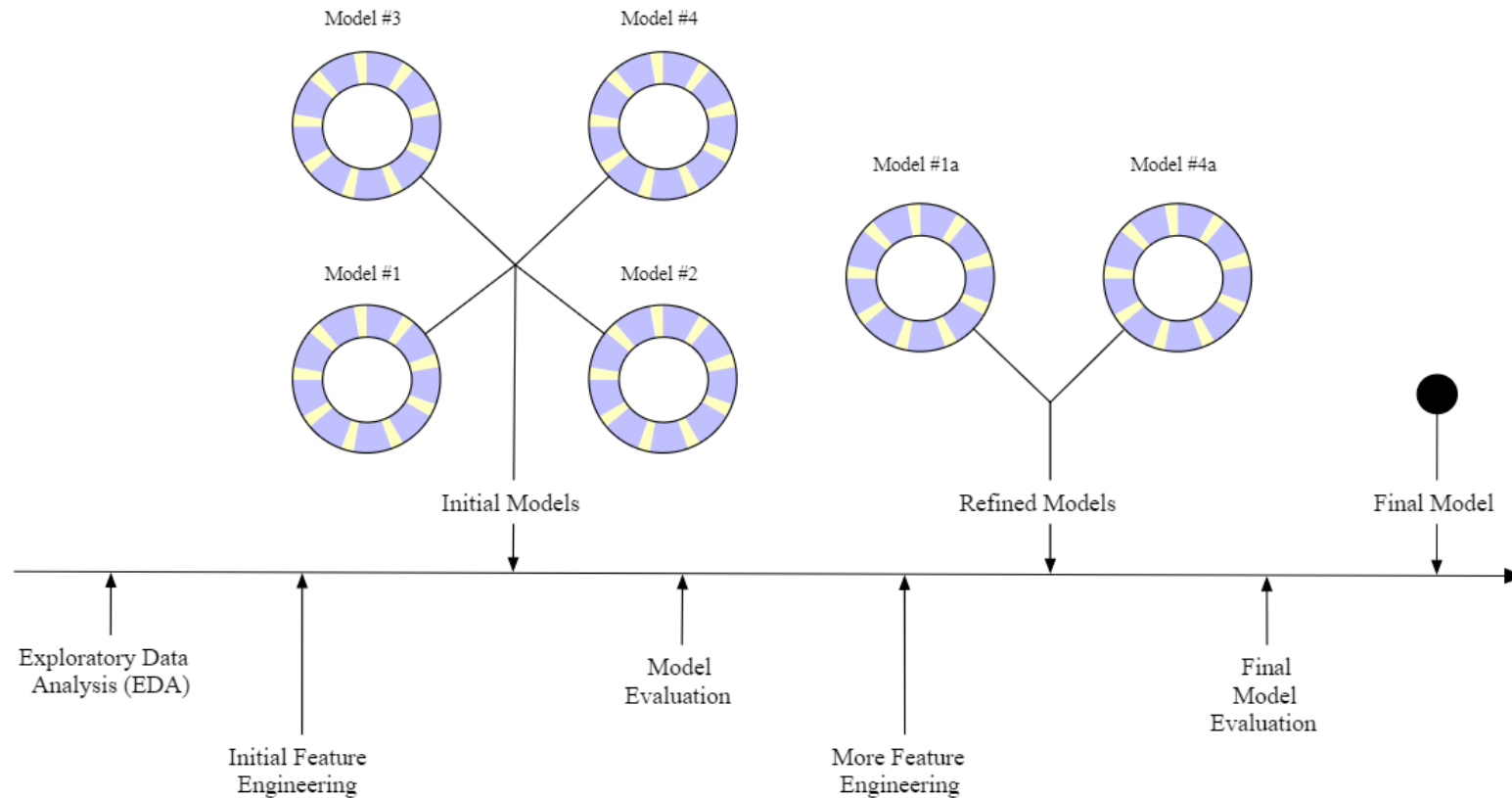
MODEL BUILDING

Typical Data Science Project



The packages shown above are designed to work together, instead of some loosely designed packages.

Typical Modeling Process



Source: *Chapter 3.3 of Tidy Modeling with R*

Issue with open-source software

Function	Package	Code
lda	MASS	predict(object)
glm	stats	predict(object, type = "response")
gbm	gbm	predict(object, type = "response", n.trees)
mda	mda	predict(object, type = "posterior")
rpart	rpart	predict(object, type = "prob")
various	RWeka	predict(object, type = "probability")
logitboost	LogitBoost	predict(object, type = "raw", nIter)
pamr.train	pamr	pamr.predict(object, type = "posterior")

Source: *Chapter 3.3 of Tidy Modeling with R*

This can be a stumbling block for users to use R to perform machine learning analysis.

Sometimes the issues can happen within the same package as well

Over here, we will take a look at this **glmnet** example shared by Max Kuhn during his **tidymodels** sharing at **Cleveland R User Group**.

glmnet is a package that allows one to fit a regularized generalized linear models. The prediction output from this package can come in various forms.

glmnet Class Predictions

```
predict(two_class_mod, newx = new_x, type = "class")
```

```
##           s0  s1  s2  
## sample_1 "a" "b" "b"  
## sample_2 "a" "b" "b"
```

glmnet Class Probabilities (Two Classes)

```
predict(two_class_mod, newx = new_x, type = "response")
```

```
##           s0  s1          s2  
## sample_1 0.5 0.5 0.5059110  
## sample_2 0.5 0.5 0.5261249
```

Now, the **predict** function returns a matrix of probability for the second level of outcome factor.

glmnet Class Probabilities (Three Classes)

```
predict(three_class_mod, newx = new_x,  
        type = "response")
```

```
## , , s0  
##  
##           a           b           c  
## sample_1 0.3333333 0.3333333 0.3333333  
## sample_2 0.3333333 0.3333333 0.3333333  
##  
## , , s1  
##  
##           a           b           c  
## sample_1 0.3333333 0.3333333 0.3333333  
## sample_2 0.3333333 0.3333333 0.3333333  
##  
## , , s2  
##  
##           a           b           c  
## sample_1 0.3727891 0.2441238 0.3830872  
## sample_2 0.3269560 0.3388499 0.3341941
```

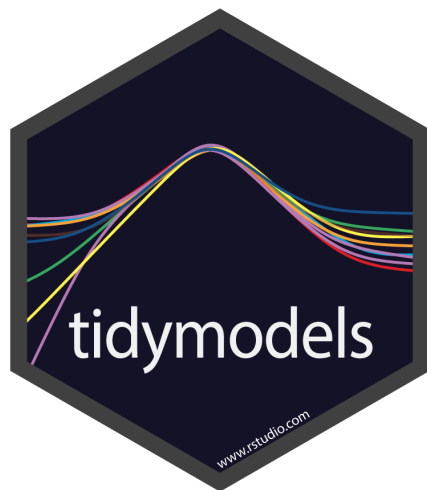
The output is no longer in matrix format. It is a 3D array format. Fainted. Often, the users would spend quite a fair bit of time in transforming the output into the required format of the next function.

Perhaps illustrating the output in such format would be better?

```
## # A tibble: 6 x 4  
##       a      b      c lambda  
##   <dbl> <dbl> <dbl> <dbl>  
## 1 0.333 0.333 0.333     1  
## 2 0.333 0.333 0.333     1  
## 3 0.333 0.333 0.333    0.1  
## 4 0.333 0.333 0.333    0.1  
## 5 0.373 0.244 0.383   0.01  
## 6 0.327 0.339 0.334   0.01
```

Tidymodels to the rescue!

The author of **caret** package (ie. Max Kuhn) felt that there should be a better approach to perform machine learning tasks.



Tidymodels is a collection of various machine learning packages, from data pre-processing to model building & model comparison. The package provides an unified interface to perform machine learning tasks.

Instead of reinventing the "wheels", the package functions as a wrapper to wrap around existing package.

Pre-Process → Train → Validate



Source: *A Gentle Introduction to tidymodels*

Above are the various key packages that assist us in our various machine learning activities.

These packages are recent project by RStudio to look into how the packages could better support users in the machine learning analysis.

The cool thing about these packages are following *tidy data* concepts.

What do you mean by tidy data?

A concept introduced by Hadley Wickham, Chief Scientist at R Studio.

Below are the definition of tidy data:

- Each variable must have its column
- Each observation must have its row
- Each value must have its cell

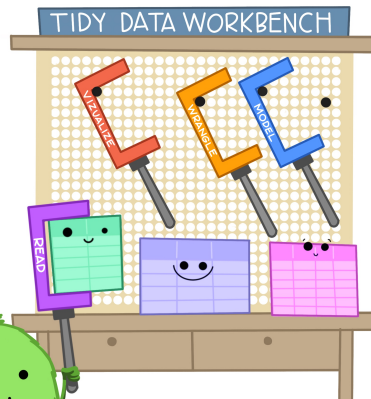
The image shows three versions of a data table with columns: country, year, cases, and population. The data rows are: Afghanistan (1999, 1666, 19999071), Afghanistan (2000, 1666, 20099360), Brazil (1999, 37737, 172006362), Brazil (2000, 80488, 174508398), China (1999, 210258, 1272912272), and China (2000, 210766, 1280425583).
1. The first diagram, labeled 'variables' below, has four vertical double-headed arrows pointing to each column header.
2. The second diagram, labeled 'observations' below, has six horizontal double-headed arrows pointing to each row.
3. The third diagram, labeled 'values' below, has circles placed inside each cell of the data rows.

country	year	cases	population
Afghanistan	1999	1666	19999071
Afghanistan	2000	1666	20099360
Brazil	1999	37737	172006362
Brazil	2000	80488	174508398
China	1999	210258	1272912272
China	2000	210766	1280425583

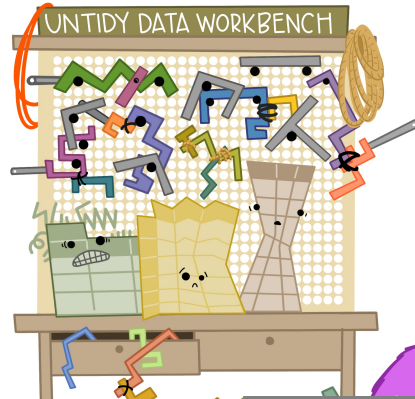
Source: *Chapter 12.2 of R for Data Science*

Hmmm, but why is this important?

When working with tidy data, we can use the **same tools** in **similar ways** for different datasets...



...but working with untidy data often means reinventing the wheel with **one-time approaches** that are **hard to iterate or reuse**.



Source: *stats-illustrations* by Allison Horst

Setting Context for the Actuarial Use Case

For the demonstration, I will be using the worker compensation insurance claim dataset from Kaggle

<https://www.kaggle.com/c/actuarial-loss-estimation/>.

Below is the snapshot of the dataset:

Show entries Search:

	ClaimNumber	DateTimeOfAccident	DateReported	Age	Gender	MaritalStatus	DependentChildren	DependentsOther
1	WC8285054	2002-04-09T07:00:00Z	2002-07-05T00:00:00Z	48	M	M	0	0
2	WC6982224	1999-01-07T11:00:00Z	1999-01-20T00:00:00Z	43	F	M	0	0
3	WC5481426	1996-03-25T00:00:00Z	1996-04-14T00:00:00Z	30	M	U	0	0

Showing 1 to 3 of 100 entries Previous 1 2 3 4 5 ... 34 Next

Data Splitting

One of the initial steps for any machine learning analysis is to split the dataset into training & testing dataset.

First, read in the clean dataset.

```
df <- read_csv("data/data_eda_actLoss_3.csv") %>%  
  drop_na()
```

Next, use **initial_split** function to create a binary split of the data into training and testing set.

```
df_split <- initial_split(df,  
                          prop = 0.6,  
                          strata = init_ult_diff)
```

training function & **testing** function are used to extract the relevant data.

```
df_train <- training(df_split)  
df_test <- testing(df_split)
```

Data Pre-processing



Source: *stats-illustrations* by Allison Horst

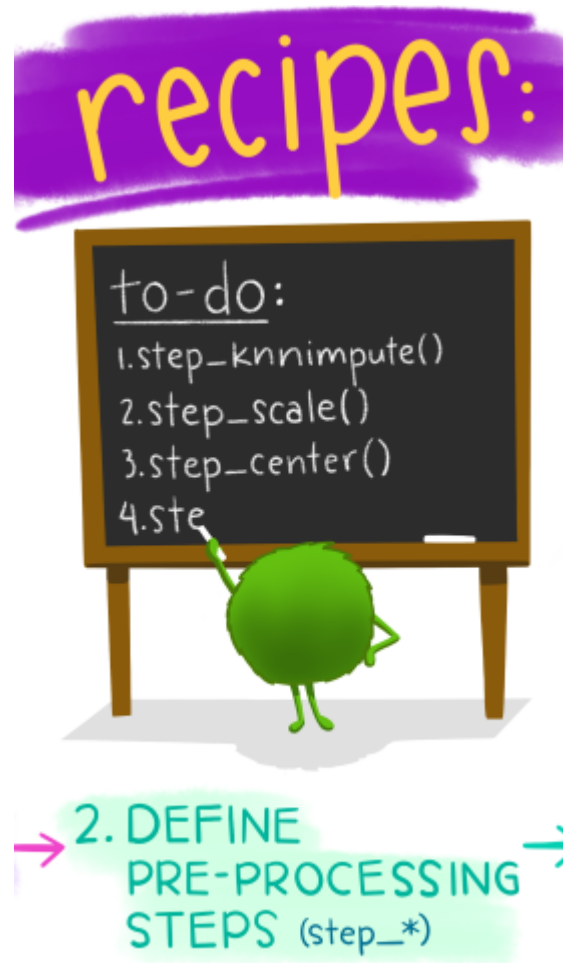
Following is the example of how the **recipe** looks like in **tidymodels**:



In line 1, I have specified the formula & dataset.

```
gen_recipe <- recipe(init_ult_diff ~ .,  
                      data = df_train)
```

Following is the example of how the **recipe** looks like in **tidymodels**:



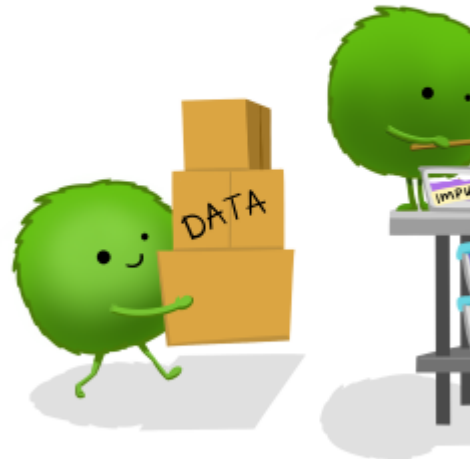
Step_date function allows users to extract year, month and day of the week from the date variable.

```
gen_recipe <- recipe(init_ult_diff ~ .,  
                      data = df_train) %>%  
  step_date(c(DateTimeOfAccident,  
              DateReported))
```

One also could perform data wrangling by using **step_mutate** function as shown under line 3 & line 4

```
gen_recipe <- recipe(init_ult_diff ~ .,  
                      data = df_train) %>%  
  step_date(c(DateTimeOfAccident,  
              DateReported)) %>%  
  step_mutate(DateTimeOfAccident_hr =  
              hour(DateTimeOfAccident),  
              DateTimeOfAccident_hr =  
              factor(DateTimeOfAccident_hr,  
                    order = TRUE))
```

Following is the example of how the **recipe** looks like in **tidymodels**:



➔ 3. PROVIDE
DATASET(S) FOR
RECIPE STEPS
prep()

Indicating the correct data type is also very important as incorrect data type would affect the model performance.

```
gen_recipe <- recipe(init_ult_diff ~ .,  
                      data = df_train) %>%  
  step_date(c(DateTimeOfAccident,  
              DateReported)) %>%  
  step_mutate(DateTimeOfAccident_hr =  
              hour(DateTimeOfAccident),  
              DateTimeOfAccident_hr =  
              factor(DateTimeOfAccident_hr,  
                    order = TRUE)) %>%  
  update_role(c(DateTimeOfAccident,  
                DateReported),  
              new_role = "id") %>%  
  prep()
```

This modeling approach has effectively allowed to modularize the different functions and chain them together.

A Peek into the Created Recipe

The code will show us the different steps we have specified under recipe step by calling the recipe object we have created.

```
gen_recipe
```

```
## Data Recipe
##
## Inputs:
##
##      role #variables
##      id      2
## outcome      1
## predictor    16
##
## Training data contained 27892 data points and no missing data.
##
## Operations:
##
## Date features from DateTimeOfAccident, DateReported [trained]
## Variable mutation for DateTimeOfAccident_hr, DateTimeOfAccident_hr, DateTimeOfAccident_year, DateReported_year [trained]
```

Checking the data type is also a very crucial step before the modeling as inappropriate variable types might affect the performance of the models.

```
gen_recipe %>%  
  summary()
```

```
## # A tibble: 26 x 4  
##   variable      type    role    source  
##   <chr>        <chr>  <chr>  <chr>  
## 1 DateTimeOfAccident date    id      original  
## 2 DateReported   date    id      original  
## 3 Age            numeric predictor original  
## 4 Gender         nominal predictor original  
## 5 MaritalStatus  nominal predictor original  
## 6 DependentChildren numeric predictor original  
## 7 DependentsOther numeric predictor original  
## 8 WeeklyWages    numeric predictor original  
## 9 PartTimeFullTime nominal predictor original  
## 10 HoursWorkedPerWeek numeric predictor original  
## # ... with 16 more rows
```

The **summary** function provides the users a quick overview of the data type.

The modularise structure allows us to effectively reuse the model components. For example, GLM model is unable to use categorical variables to fit the model.

So, to resolve this, we can add on additional data preprocessing step.

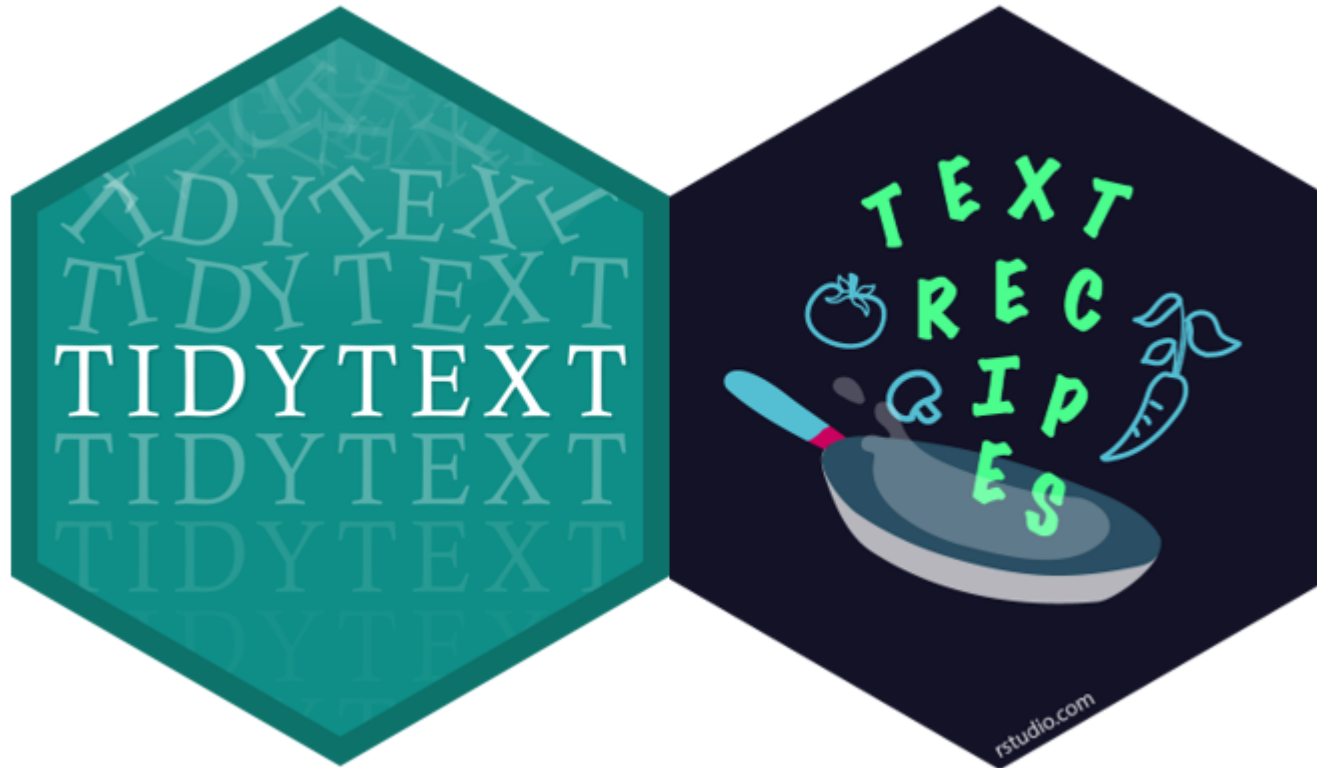
```
glmnet_recipe <- gen_recipe %>%  
  step_dummy(all_nominal())
```

The code above indicates that one-hot encoding method to be applied on all the categorical variables .

Wait! Don't throw away the text fields

Remember there is a claim description field within the data?

This is where **tidytext** package and **textrecipe** comes very handy.



tidytext

To extract the text, I will first split the words into tokens by using **unnest_token**.

```
tidy_clm_unigram <- data_1 %>%  
  unnest_tokens(word, ClaimDescription,  
                token = "ngrams",  
                n = 1)
```

I will also remove the stopwords (eg. above, onto, and) from the tokens since they are not meaningful in the analysis.

```
tidy_clm_unigram <- data_1 %>%  
  unnest_tokens(word, ClaimDescription,  
                token = "ngrams",  
                n = 1) %>%  
  anti_join(get_stopwords())
```

Once the words are tokenized, we can perform frequency count on the words to understand which are the words tend to appear more frequent than the rest.

```
tidy_clm_unigram %>%  
  count(word, sort = TRUE)
```

```
## # A tibble: 3,449 x 2  
##   word      n  
##   <chr>   <int>  
## 1 right   19839  
## 2 left    18470  
## 3 back    13438  
## 4 strain  12124  
## 5 lower    8170  
## 6 finger   7884  
## 7 hand     7145  
## 8 struck   6868  
## 9 lifting  6774  
## 10 eye     5155  
## # ... with 3,439 more rows
```

After that, we can create indicators for the more frequently appeared words to see whether the model performance would improve by including these features.

Once they are extracted, we can visualize the extracted texts by using word cloud.

```
cleaned_clm_unigram %>%
  filter(n > 2000) %>%
  ggplot(aes(label = word,
              size = n,
              color = n)) +
  geom_text_wordcloud() +
  scale_size_area(max_size = 20) +
  theme_minimal()
```



textrecipe

Alternatively, we can perform text mining through specifying the steps in the recipe.

Same as the previous recipe created, after specifying the 'recipe' for the machine learning model, we will specify what are the pre-processing steps the model should perform.

```
ranger_recipe_clmdesc <-  
  recipe(formula = init_ult_diff ~ .,  
          data = df_wClmDesc_train) %>%  
    step_tokenize(ClaimDescription) %>%  
    step_stopwords(ClaimDescription) %>%  
    step_tokenfilter(ClaimDescription,  
                     max_tokens = 20) %>%  
    step_tfidf(ClaimDescription)
```

- This approach is consistent with recipe to "prepare" the text data for modeling
- It is easier to understand and clear at first glance what are we "preparing"

More explanation on the different text mining techniques and how they work, do check this [book](#) or this [package](#).

Model Selection - Model X, I choose you!



Back to Actuaries' Most Beloved Model - GLM

Convention Approach

```
x <- df_train %>%  
  dplyr::select(-init_ult_diff) %>%  
  data.matrix()  
  
y <- df_train$init_ult_diff  
  
glmnet(x,  
  y,  
  family = "gaussian",  
  nlambda = 10,  
  alpha = 0.5)
```

Tidymodels Approach

```
glmnet_spec <-  
  linear_reg(penalty = tune(),  
             mixture = tune()) %>%  
  set_mode("regression") %>%  
  set_engine("glmnet", family = "gaussian")
```

parsnip package provides users a more unified model interface without sacrificing the flexibility.

parsnip package do support a wide range of machine learning models.

Random Forest

```
ranger_spec <-  
  rand_forest(mtry = tune(),  
              min_n = tune(),  
              trees = 50) %>%  
  set_mode("regression") %>%  
  set_engine("ranger",  
             importance = "impurity")
```

XGBoost

```
xgboost_spec <-  
  boost_tree(trees = tune(),  
             min_n = tune(),  
             tree_depth = tune(),  
             learn_rate = tune(),  
             loss_reduction = tune(),  
             sample_size = tune()) %>%  
  set_mode("regression") %>%  
  set_engine("xgboost")
```

Linear Regression

```
lm_spec <- linear_reg() %>%  
  set_mode("regression") %>%  
  set_engine("lm")
```

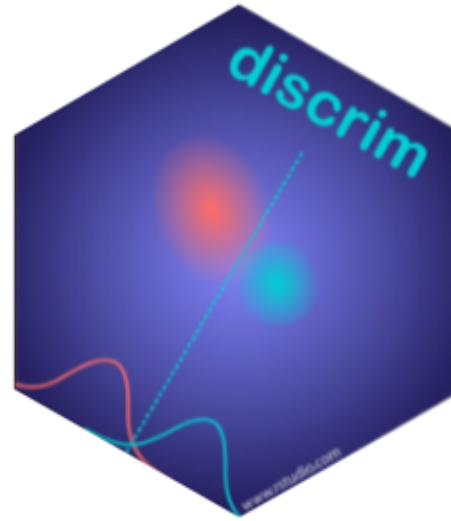
MARS

```
earth_spec <-  
  mars(num_terms = tune(),  
        prod_degree = tune(),  
        prune_method = "none") %>%  
  set_mode("regression") %>%  
  set_engine("earth")
```

KNN

```
kknn_spec <-  
  nearest_neighbor(neighbors = tune(),  
                  weight_func = tune()) %>%  
  set_mode("regression") %>%  
  set_engine("kknn")
```

There are other tidymodels packages support different machine learning models...

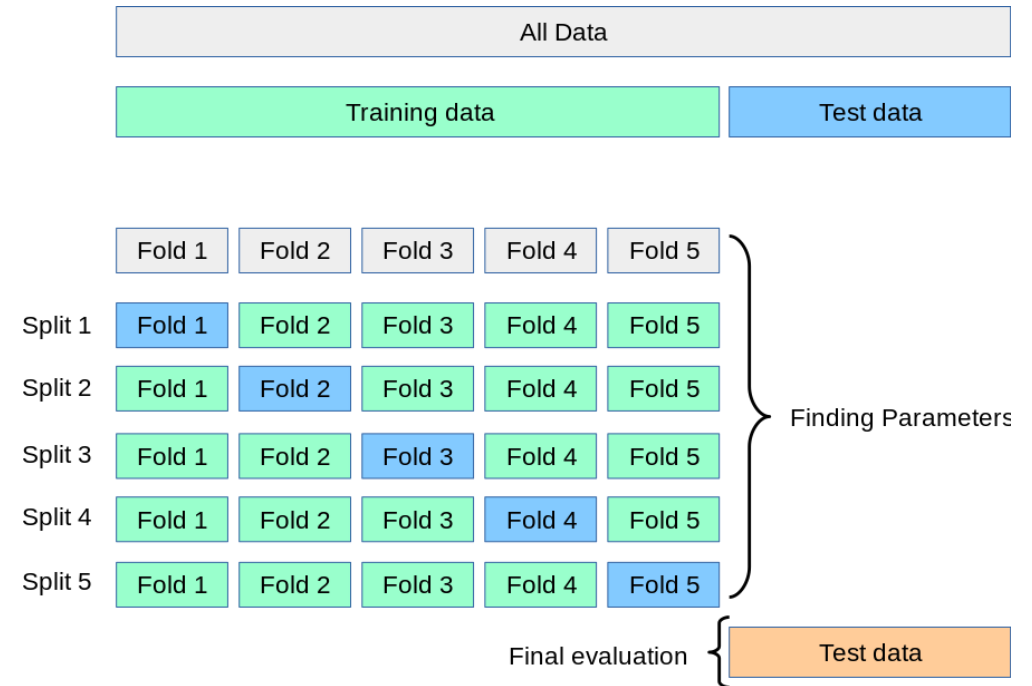


Model Tuning

Cross validation is commonly used to:

- Find the best set of parameters that would give us best model performance
- Prevent models from overfitting

Over here, I will be using k-fold validation to perform model tuning.



Source: *Section 3.1 of Scikit-learn*

To do so, we will create the dataset for k-fold cross validation by using **vfold_cv** function as shown below.

```
df_folds <- vfold_cv(df_train, strata = init_ult_diff)
df_folds
```

And yes we are almost there!

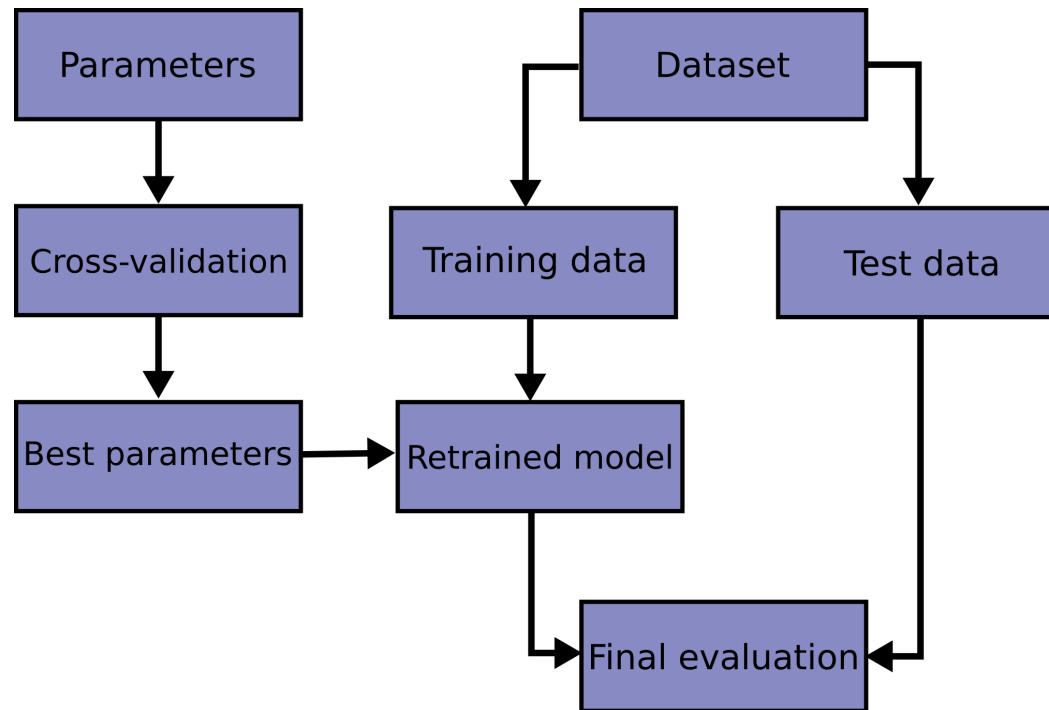
Over here, **grid search** is used to find the best parameters fit a given model.

```
ranger_tune <-  
  tune_grid(ranger_workflow,  
            resamples = df_folds,  
            grid = 5)
```

```
ranger_fit <- ranger_workflow %>%  
  finalize_workflow(select_best(ranger_tune)) %>%  
  last_fit(df_split)
```

Once the model is tuned, **select_best** function is used to select the best parameters. Then, finalise the workflow & run the fitted model on testing data by using **last_fit** function.

If we were to visualize the flow, the modeling steps mentioned above would look something as following....



Now, we can "chain" the different components as a workflow

Once the necessary parameters are setup, we will join the different steps together to form a workflow.

Conventional Approach

```
x <- df_train %>%  
  dplyr::select(-init_ult_diff) %>%  
  data.matrix()  
  
y <- df_train$init_ult_diff  
  
glmnet(x,  
  y,  
  family = "gaussian",  
  nlambda = 10,  
  alpha = 0.5)
```

tidymodels Approach

```
glmnet_workflow <-  
  workflow() %>%  
  add_recipe(glmnet_recipe) %>%  
  add_model(glmnet_spec)
```

Need another workflow? Not an issue, sir

```
glmnet_workflow_ult <-  
  glmnet_workflow %>%  
  update_formula(UltimateIncurredClaimCost ~ .)
```

The created workflow can only be visualised by calling the workflow item.

```
glmnet_workflow
```

```
## == Workflow =====  
## Preprocessor: Recipe  
## Model: linear_reg()  
##  
## -- Preprocessor -----  
## 3 Recipe Steps  
##  
## * step_date()  
## * step_mutate()  
## * step_dummy()  
##  
## -- Model -----  
## Linear Regression Model Specification (regression)  
##  
## Main Arguments:  
##   penalty = tune()  
##   mixture = tune()  
##  
## Computational engine: glmnet
```

Such modularized modeling method allows us to reuse the different machine learning parts through the analysis.

Confused?? Worry not

usemodels package can be used to generate all the necessary modeling templates for the users.

```
use_ranger(formula, data)
```

```
ranger_recipe <-  
  recipe(formula = init_ult_diff ~ ., data = df_train) %>%  
  step_string2factor(one_of(MaritalStatus, PartTimeFullTime, injury_body, injury_side,  
    injury_item, injury_cause, injury_type))  
  
ranger_spec <-  
  rand_forest(mtry = tune(), min_n = tune(), trees = 1000) %>%  
  set_mode("regression") %>%  
  set_engine("ranger")  
  
ranger_workflow <-  
  workflow() %>%  
  add_recipe(ranger_recipe) %>%  
  add_model(ranger_spec)  
  
set.seed(93447)  
ranger_tune <-  
  tune_grid(ranger_workflow, resamples = stop("add your rsample object"), grid = stop("add number of candidate points"))
```

Model Comparison

To facilitate the model comparison, **yardstick** is being used to compare the different models.



Same as other packages under **tidymodels**, various model performance metrics under **yardstick** also have same interface, which allows us to loop through the various syntax to calculate the model performance.

First, I have defined the different measurements I need.

I will be using *root mean square error*, *R squared* and *mean absolute scaled error*.

Root Mean Square Error

```
rmse(data, truth, estimate, na_rm = TRUE, ...)
```

R Squared

```
rsq(data, truth, estimate, na_rm = TRUE, ...)
```

Mean Absolute Scaled Error

```
mase(data, truth, estimate, m = 1L, mae_train = NULL, na_rm = TRUE, ...)
```

As shown above, the metric interfaces look very similar. This allows to calculate the model performance by looping through the different metrics.

```
model_metrics <- metric_set(rmse, rsq, mase)
```

```
ranger_pred <- ranger_fit %>%  
  collect_predictions()  
  
ranger_metric <- model_metrics(ranger_pred,  
                               truth = init_ult_diff,  
                               estimate = .pred) %>%  
  mutate(model = "ranger") %>%  
  pivot_wider(names_from = .metric,  
              values_from = .estimate)
```

Next, I will join these model results into a tibble table so that I can compare how does the different models perform under different scenarios.

Showdown by Different Models

Following are the performance metrics of different models:

```
## # A tibble: 6 x 5
##   .estimator model    rmse   rsq  mase
##   <chr>      <chr>  <dbl> <dbl> <dbl>
## 1 standard  ranger  1537. 0.457 0.490
## 2 standard  xgboost 1577. 0.441 0.511
## 3 standard  glmnet  1656. 0.368 0.564
## 4 standard  lm      1695. 0.338 0.569
## 5 standard  earth   1708. 0.328 0.593
## 6 standard  kknn    1758. 0.288 0.581
```

In general, the different model performance results are quite consistent with one another.

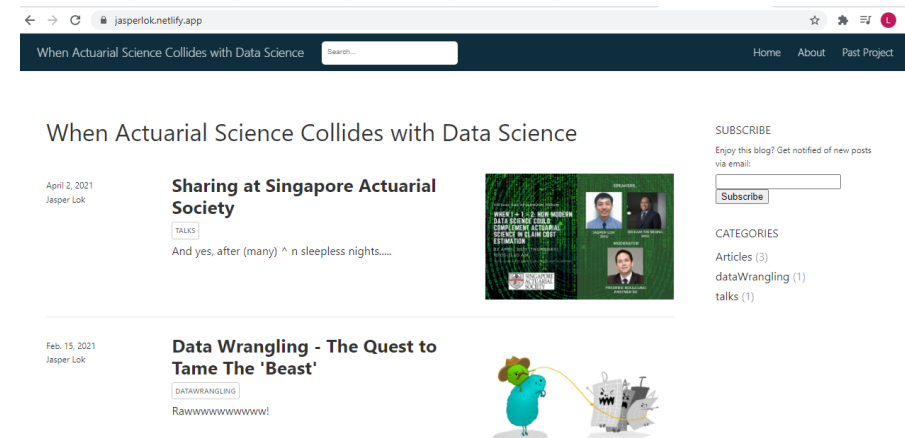
Out of all the models, random forest has the highest accuracy.

This is just a very small subset of the entire research



The full research paper and R code are published on my Github https://github.com/jasperlok/SMU_Capstone.

This deck of interactive slides will be shared on my data science blog, [When Actuarial Science Collides with Data Science](#).



FAQ

Thank you!

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Blog: <https://jasperlok.netlify.app/>

Appendix

Link to the relevant R packages

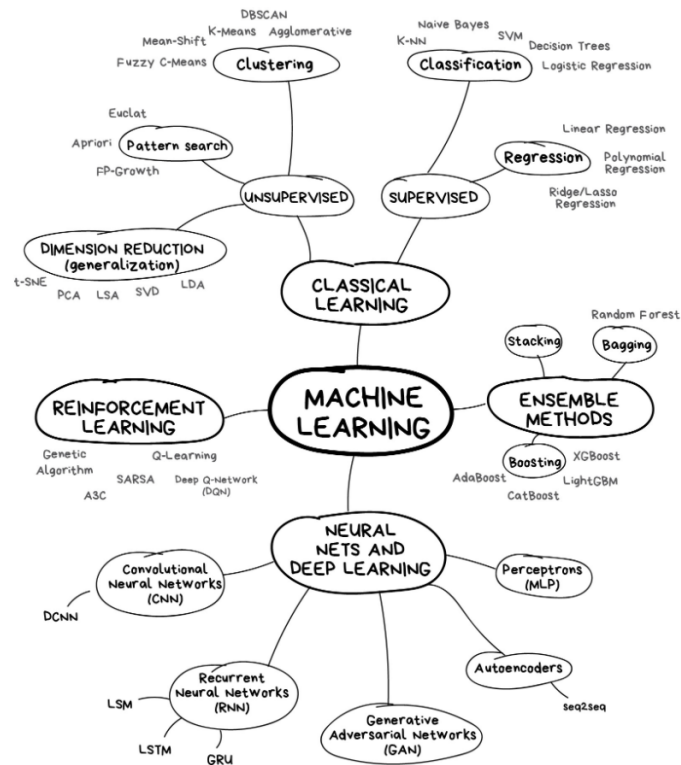
Tidyverse <https://www.tidyverse.org/>

Tidymodels <https://www.tidymodels.org/>

Variable Importance Plot <https://koalaverse.github.io/vip/index.html>

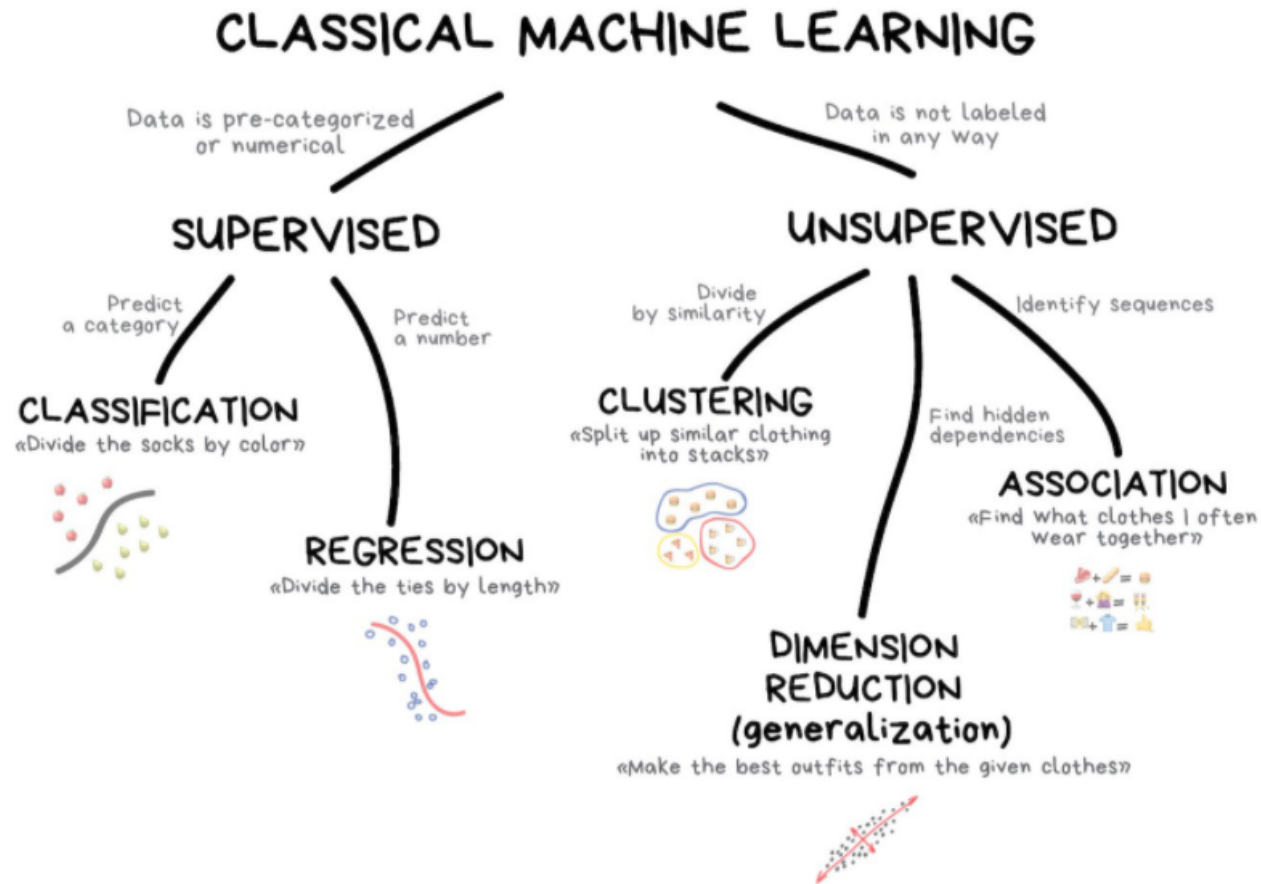
Partial Dependence Plot <https://bgreenwell.github.io/pdp/index.html>

Overview of Machine Learning



Source: *xkcd* blog

Overview of Classical Machine Learning



Further reading

R for Data Science

<https://r4ds.had.co.nz/>

Tidymodeling with R

<https://www.tmwr.org/>

Feature Engineering

<https://bookdown.org/max/FES/>

Text Mining

<https://www.tidytextmining.com/>

Interpretable Machine Learning

<https://christophm.github.io/interpretable-ml-book/>