HYPERPARAMETER TUNING IN R

Parameters vs hyperparameters

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About me

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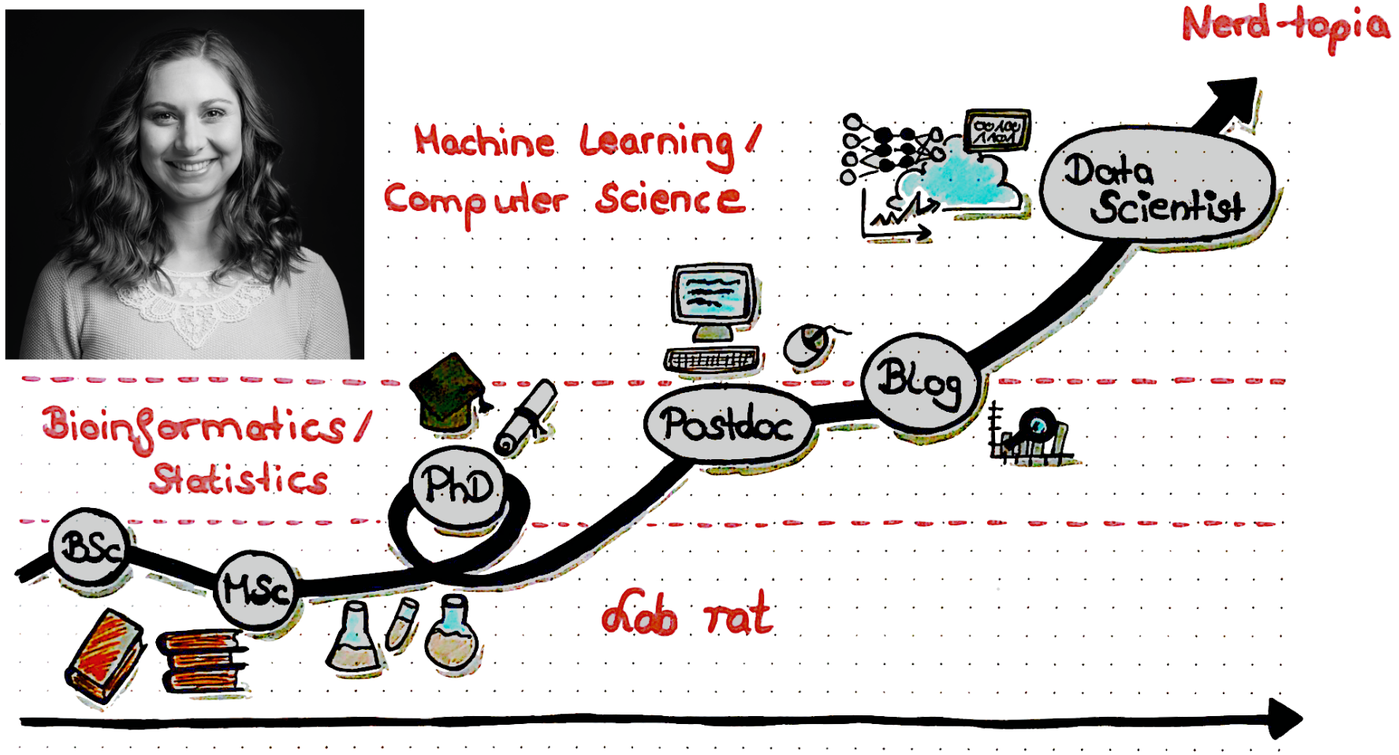


"Hyper"parameters vs model parameters

- Let's look at an example dataset:

```
head(breast_cancer_data)

# A tibble: 6 x 11
  diagnosis concavity_mean symmetry_mean fractal_dimension_… perimeter_se smooth
  <chr>             <dbl>          <dbl>                <dbl>        <dbl>
1 M                 0.300          0.242               0.0787         8.59
2 M                 0.0869         0.181               0.0567         3.40
3 M                 0.197          0.207               0.0600         4.58
4 M                 0.241          0.260               0.0974         3.44
5 M                 0.198          0.181               0.0588         5.44
6 M                 0.158          0.209               0.0761         2.22
```

- And build a simple **linear model**.

Let's start simple: Model parameters in a linear model

```
# Create linear model
linear_model <- lm(perimeter_worst ~ fractal_dimension_mean,
                   data = breast_cancer_data)

# Get fitted model parameters
summary(linear_model)

# Get residuals
resid(linear_model)

    Min      1Q  Median      3Q     Max
-50.094 -24.859  -7.705  22.209  89.919

# Get coefficients
linear_model$coefficients

                       Estimate Std. Error t value Pr(>|t|)
(Intercept)              167.60      25.91   6.469  3.9e-09 ***
fractal_dimension_mean  -926.39     392.86  -2.358   0.0204 *
```

Let's start simple: Model parameters in a linear model

- Model **parameters** are being fit (i.e. found) during training.
- They are the **result** of model fitting or training.
- In a linear model, we want to find the **coefficients**.

```
> linear_model$coefficients

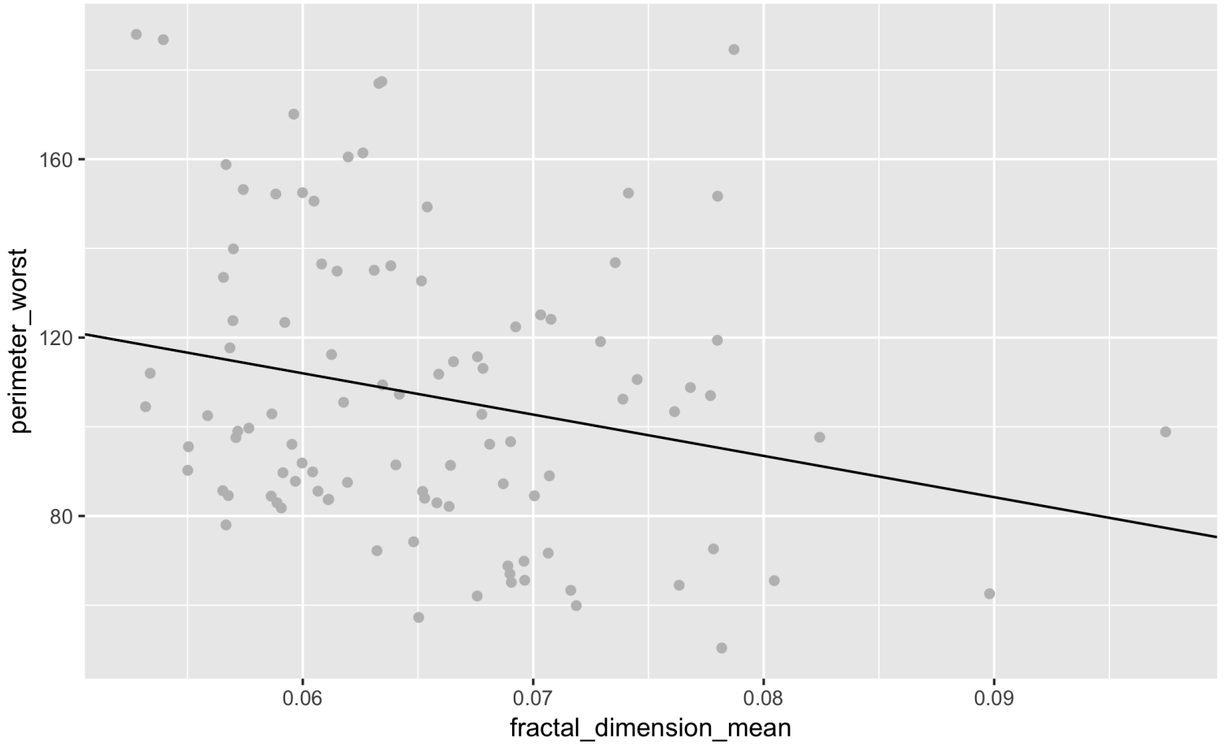
           (Intercept) fractal_dimension_mean
              167.5972               -926.3866
```

- We can think of them as the **slope** and the **y-intercept** of our model.

Coefficients in a linear model

```
ggp <- ggplot(data = breast_cancer_data,
              aes(x = fractal_dimension_mean, y = perimeter_worst)) +
       geom_point(color = "grey")

ggp + geom_abline(slope = linear_model$coefficients[2],
                  intercept = linear_model$coefficients[1])
```



Model parameters vs hyperparameters in a linear model

- *Remember*: model **parameters** are being fit (i.e. found) during training; they are the **result** of model fitting or training.
- **Hyperparameters** are being set before training.
- They specify **HOW** the training is supposed to happen.

```
args(lm)
formals(lm)
help(lm)
?lm

linear_model <- lm(perimeter_worst ~ fractal_dimension_mean,
                   data = breast_cancer_data,
                   method = "qr")
```

Parameters vs hyperparameters in machine learning

In our **linear model**:

- Coefficients were **found** during fitting.
- `method` was an option to set **before** fitting.

In **machine learning** we might have:

- Weights and biases of neural nets that are optimized during training => model **parameters**.
- Options like learning rate, weight decay and number of trees in a Random Forest model that can be tweaked => **hyperparameters**.

Why tune hyperparameters?

- Fantasy football players ~ **Hyperparameters**
- Football players' positions ~ Hyperparameter **values**

- Finding the best combination of players and positions ~ Finding the best **combination** of hyperparameters

HYPERPARAMETER TUNING IN R

Let's practice!

HYPERPARAMETER TUNING IN R

Machine Learning with caret - the Basics

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Machine Learning with caret - splitting data

- **Splitting** into training and test data:

```
# Load caret and set seed
library(caret)
set.seed(42)

# Create partition index
index <- createDataPartition(breast_cancer_data$diagnosis, p = .70,
                             list = FALSE)

# Subset `breast_cancer_data` with index
bc_train_data <- breast_cancer_data[index, ]
bc_test_data  <- breast_cancer_data[-index, ]
```

- Training set with enough **power**.
- **Representative** test set.

Train a machine learning model with caret

- Set up **cross-validation**:

```
library(caret)
library(tictoc)

# Repeated CV.
fitControl <- trainControl(method = "repeatedcv",
                           number = 3,
                           repeats = 5)
```

- **Train** a Random Forest model:

```
tic()
set.seed(42)
rf_model <- train(diagnosis ~ .,
                  data = bc_train_data,
                  method = "rf",
                  trControl = fitControl,
                  verbose = FALSE)
toc()

1.431 sec elapsed
```

Automatic hyperparameter tuning in caret

```
rf_model

Random Forest

80 samples
10 predictors
 2 classes: 'B', 'M'

No pre-processing
Resampling: Cross-Validated (3 fold, repeated 5 times)
Summary of sample sizes: 54, 54, 52, 54, 53, 53, ...
Resampling results across tuning parameters:

  mtry  Accuracy   Kappa
   2    0.9006783  0.8015924
   6    0.9126645  0.8253289
  10    0.8999389  0.7999386

Accuracy was used to select the optimal model using the largest value.
The final value used for the model was mtry = 6.
```

HYPERPARAMETER TUNING IN R

Let's start modeling!

HYPERPARAMETER TUNING IN R

Hyperparameter tuning with caret

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Automatic hyperparameter tuning in caret

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rf_model

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```

Hyperparameters are specific to model algorithms

- `modelLookup(model)`
- https://topepo.github.io/caret/available-models.html

Hyperparameters in Support Vector Machines (SVM)

```
> library(caret)
> library(tictoc)

> fitControl <- trainControl(method = "repeatedcv",
                             number = 3,
                             repeats = 5)
tic()
set.seed(42)
svm_model <- train(diagnosis ~ .,
                   data = bc_train_data,
                   method = "svmPoly",
                   trControl = fitControl,
                   verbose= FALSE)
toc()

3.836 sec elapsed
```

Hyperparameters in Support Vector Machines (SVM)

```
svm_model

Support Vector Machines with Polynomial Kernel

...

Resampling results across tuning parameters:

  degree  scale  C     Accuracy   Kappa
 ...
  1       0.100  1.00  0.9104803  0.8211459
 ...

Accuracy was used to select the optimal model using the largest value.
The final values used for the model were degree = 1, scale = 0.1 and C = 1.
```

Defining hyperparameters for automatic tuning

- **tuneLength**

```
tic()
set.seed(42)
svm_model_2 <- train(diagnosis ~ .,
                     data = bc_train_data,
                     method = "svmPoly",
                     trControl = fitControl,
                     verbose = FALSE,
                     tuneLength = 5)
toc()

7.458 sec elapsed

svm_model_2

...

Accuracy was used to select the optimal model using the largest value.
The final values used for the model were degree = 1, scale = 1 and C = 1.
```

Manual hyperparameter tuning in caret

- **tuneGrid + expand.grid**

```
library(caret)
library(tictoc)

hyperparams <- expand.grid(degree = 4,
                           scale = 1,
                           C = 1)

tic()
set.seed(42)
svm_model_3 <- train(diagnosis ~ .,
                     data = bc_train_data,
                     method = "svmPoly",
                     trControl = fitControl,
                     tuneGrid = hyperparams,
                     verbose = FALSE)
toc()

0.691 sec elapsed
```

Manual hyperparameter tuning in caret

```
svm_model_3

Support Vector Machines with Polynomial Kernel

...

  Accuracy   Kappa
  0.7772947  0.554812

Tuning parameter 'degree' was held constant at a value of 4
Tuning parameter 'scale' was held constant at a value of 1
Tuning parameter 'C' was
 held constant at a value of 1
```

HYPERPARAMETER TUNING IN R

It's your turn!