

Parameters vs hyperparameters

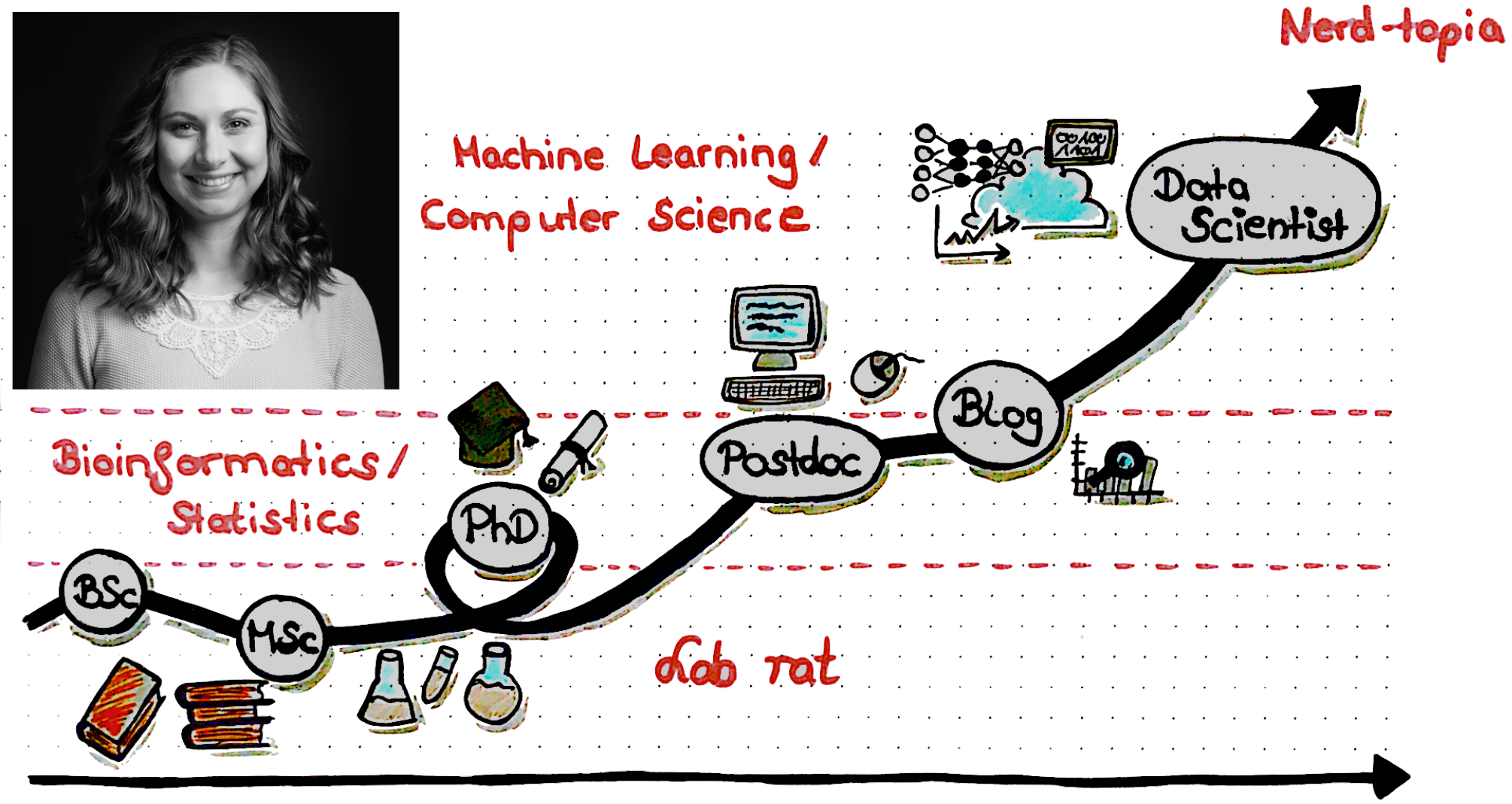
HYPERPARAMETER TUNING IN R



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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 smoothness_se
  <chr>          <dbl>          <dbl>          <dbl>          <dbl>          <dbl>
1 M           0.300          0.242          0.0787          8.59          0.00640
2 M           0.0869         0.181          0.0567          3.40          0.00522
3 M           0.197          0.207          0.0600          4.58          0.00615
4 M           0.241          0.260          0.0974          3.44          0.00911
```

- 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 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**.

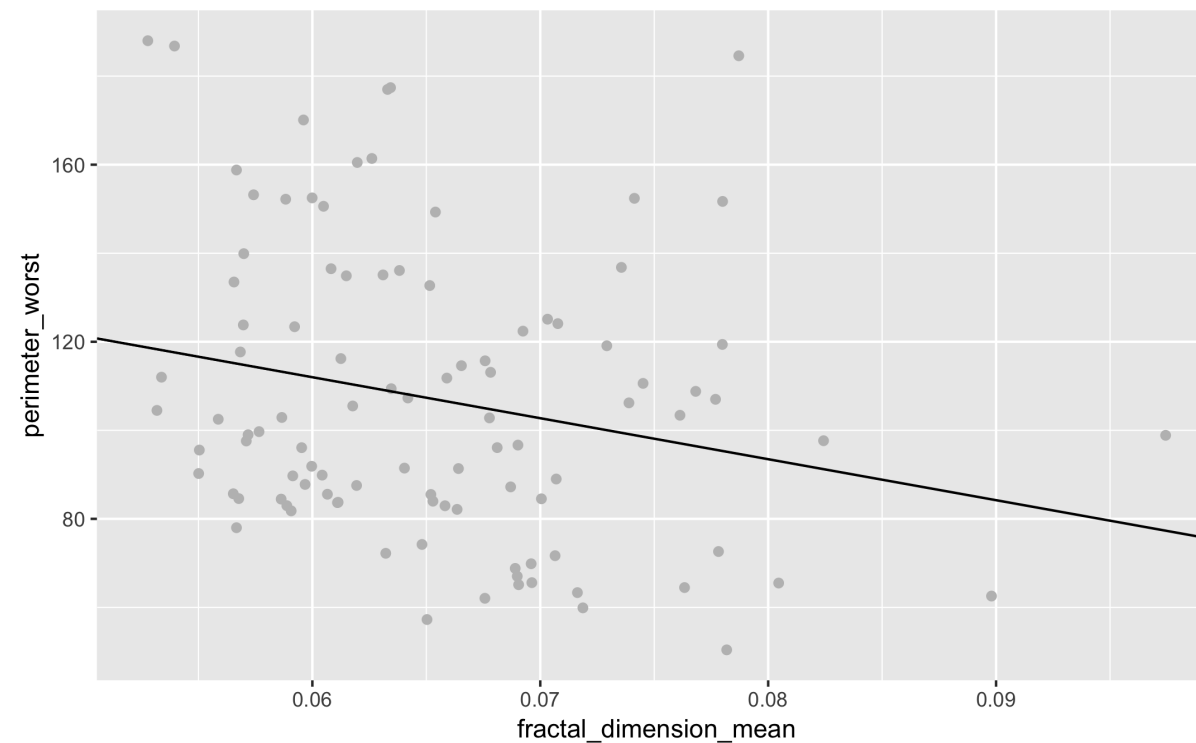
```
linear_model$coefficients
```

```
(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)
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



Let's practice!
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Recap of machine learning basics

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Machine learning with caret - splitting 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)
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

```
Random Forest
```

```
...
```

```
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.
```

Let's start modeling!

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Hyperparameter tuning with caret

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Automatic hyperparameter tuning in caret

```
Random Forest
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```
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Hyperparameters are specific to model algorithms

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

6 Available Models

The models below are available in `train`. The code behind these protocols can be obtained using the function `getModelInfo` or by going to the [github repository](https://topepo.github.io/caret/available-models.html).

Show entries

Search:

Model <input type="checkbox"/>	<i>method</i> Value <input type="checkbox"/>	Type <input type="checkbox"/>	Libraries <input type="checkbox"/>	Tuning Parameters
AdaBoost Classification Trees	adaboost	Classification	fastAdaboost	nIter, method
AdaBoost.M1	AdaBoost.M1	Classification	adabag, plyr	mfinal, maxdepth, coflearn
Adaptive Mixture Discriminant Analysis	amdai	Classification	adaptDA	model
Adaptive- Network Based				

Hyperparameters in Support Vector Machines (SVM)

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

tic()
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

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`

```
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
```

It's your turn!

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