## Building meaningful machine learning models for disease prediction

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#### About me

since 2015 Bioinformatics Postdoc Next Generation Sequencing autoinflammatory diseases & innate immunity



2011 - 2015 PhD in Biology
Is the immune system of plants required to adapt to flowering time change?

2005 - 2011 BSc and MSc of Science in Biology evolutionary genetics, immune memory in Drosophila

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## What makes a model meaningful?

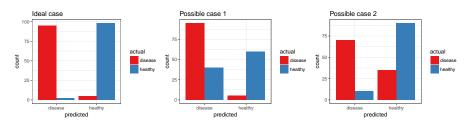
## What makes a model meaningful?

- creating ML models is relatively easy
- creating good or meaningful models is hard

#### Meaningful models

- are generalizable
- answer the question(s) posed...
- ... with sufficient accuracy to be trustworthy

## Accuracy depends on the problem!





## ML in disease modeling

- tools that can interpret "big medical data"
- and provide fast, accurate and actionable information
- for precision or personalized medicine

#### Examples:

- computer-aided diagnosis of breast cancer from mammograms<sup>1</sup>
- identifying gene defects with facial recognition software<sup>2</sup>
- identifying signatures of Brain Cancer from MRSI<sup>3</sup>
- ... and many more ...



<sup>&</sup>lt;sup>1</sup>Doi 2007.

<sup>&</sup>lt;sup>2</sup>Levenson 2014.

<sup>&</sup>lt;sup>3</sup>Sadja 2006.

# Can we trust a model?



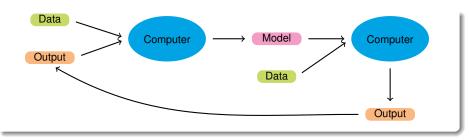
- most ML algorithms model high-degree interactions between variables
- we often don't know WHY ML models make decisions
- inherent problem with ML models: they are hard (or impossible) to interpret
- therefore, it is crucial that our models are meaningful

Image source: Pixabay

## A quick recap of ML basics

## Machine learning

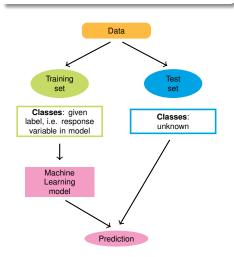
- artificial intelligence (AI)
- data-driven
- algorithms learn by being trained on observed data...
- ... and predict unknown data
- ML concepts are not new, but the increase in computational capacity has made them more accessible

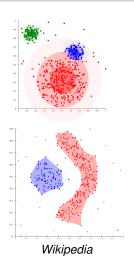


## Supervised vs Unsupervised algorithms

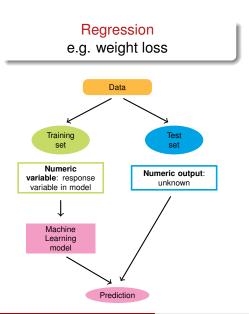
#### Supervised

#### Unsupervised





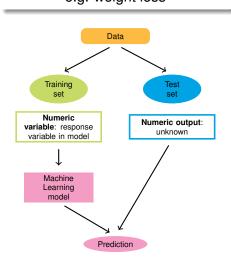
## Classification vs Regression

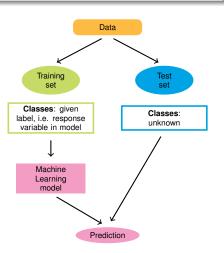


## Classification vs Regression

## Regression e.g. weight loss

### Classification e.g. healthy vs disease

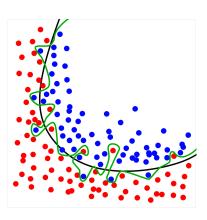




#### **Features**

- Features are the variables used for model training.
- Using the right features is crucial.

- More is not necessarily better (overfitting)!
- feature selection
- feature extraction/ engineering



Wikipedia



## Hyper Parameter Tuning

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## Take home messages:

• ...

#### How to build ML models in R

## Session setup

Breast Cancer Wisconsin Dataset<sup>4</sup>





- caret<sup>5</sup>
- h2o<sup>6</sup>

### Code will be available on my website and on Github

Friday, 31st March 2017

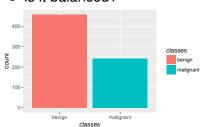
<sup>&</sup>lt;sup>4</sup>W. H. Wolberg and O. L. Mangasarian (1990). "Multisurface method of pattern separation for medical diagnosis applied to breast cytology." In: *Proceedings of the National Academy of Sciences* 87.23, pp. 9193–9196.

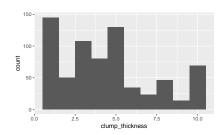
<sup>&</sup>lt;sup>5</sup>M. Kuhn et al. (2016). *caret: Classification and Regression Training*. R package version 6.0-71.

<sup>&</sup>lt;sup>6</sup>H2O.ai (2017). *h2o: R Interface for H2O*. . R package version 3.10.3.6.

#### Response variable

Is it balanced?

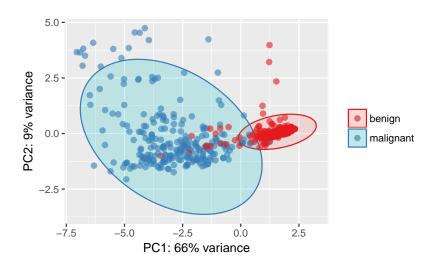




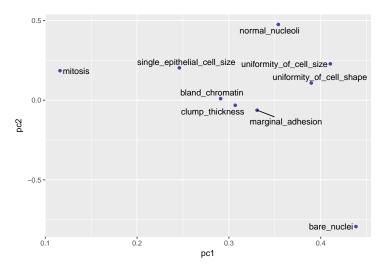
#### Missing data

- Is there missing data?
- Can we afford to loose data points with missing values?
- Or do we use imputation (and introduce additional uncertainty)?

#### Principal Component Analysis (PCA)

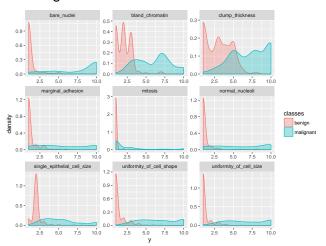


#### Principal Component Analysis (PCA)

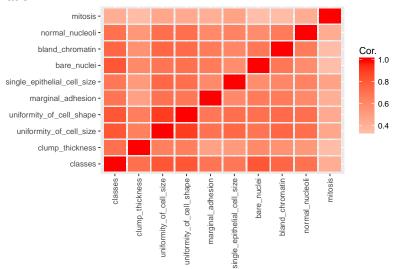


#### **Features**

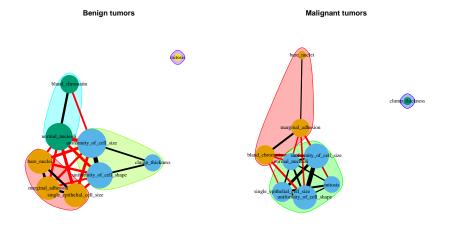
- factors or numeric
- pre-processing



#### Correlation



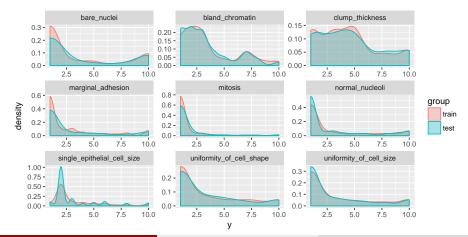
#### Correlation graphs



## Training, validation and test data

We need to split the data into training and test sets - ideally stratified by response class.

#### **Density distribution**



## Model examples

#### Regression with Linear Models

- e.g. Generalized Linear Models
- example with caret

#### Tree-based classification

- Random Forest or Gradient boosting trees
- with caret

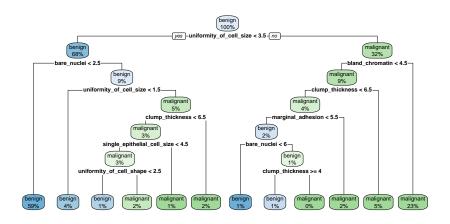
#### Hyper-parameter tuning

- Grid Search
- with h2o

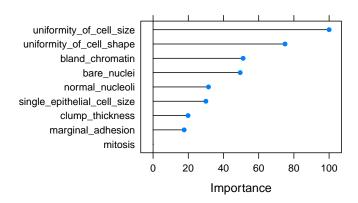
### Classification with tree-based models

#### **Decision trees**

e.g. Random Forest and gradient boosting trees



## Feature importance



## Evaluating model performance

Never use the same data

for evaluation that you used

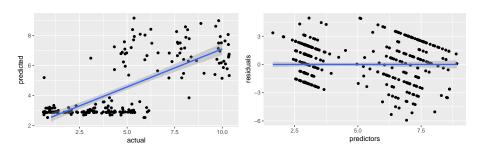
for training!

### Predictions on test data

#### Regression

• RMSE: 1.97

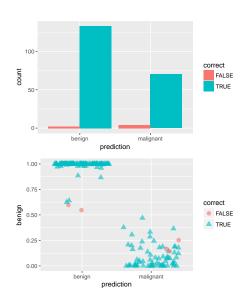
• R<sup>2</sup>: 0.50



#### Predictions on test data

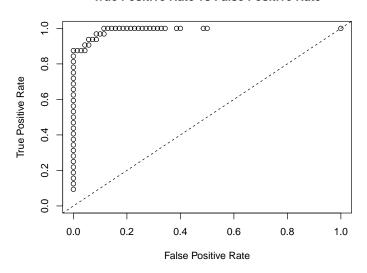
#### Classification

```
## Confusion Matrix and Statistics
##
##
              Reference
## Prediction benign malignant
##
     benign
                  133
##
     malignant
                             70
##
##
                  Accuracy: 0.9713
##
                    95% CI: (0.9386, 0.9894)
       No Information Rate: 0.6555
##
##
       P-Value [Acc > NIR] : <2e-16
##
##
                     Kappa: 0.9369
    Mcnemar's Test P-Value: 0.6831
##
##
##
               Sensitivity: 0.9708
               Specificity: 0.9722
##
##
            Pos Pred Value : 0.9852
            Neg Pred Value: 0.9459
##
##
                Prevalence: 0.6555
##
            Detection Rate: 0.6364
      Detection Prevalence: 0.6459
##
##
         Balanced Accuracy: 0.9715
##
          'Positive' Class : benign
##
##
```

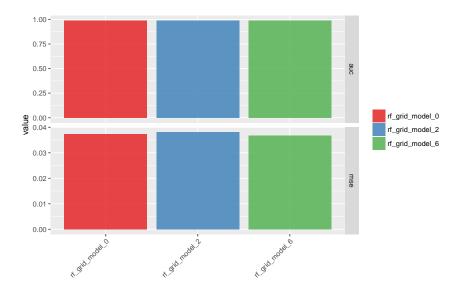


## Area Under the Curve (AUC

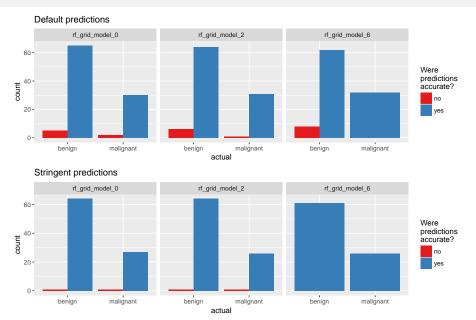
#### True Positive Rate vs False Positive Rate



## AUC and mean squared error (MSE)



### Predictions on test data



## Take home messages:

• ...

#### Outlook

- 'big data' needs to be big!
- for really meaningful models, data needs to be shared
- the more data, the more accurate and generalizable the models will be
- issues: privacy, platform, quality standards
- ML could make health care more cost-effective by reducing the energy required for interpretation

## Thank you for your attention!

## Questions?

Slides and code will be available on Github: https://github.com/ShirinG/Webinar\_ML\_for\_disease

Code will also be on my website: https://shiring.github.io

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