About me

prof. dr. Sven Degroeve

Master in Informatics, PhD in Computer Sciences

Computational Omics and Systems Biology Group (CompOmics, prof. dr. Lennart Martens)

Vakgroep Biomoleculaire Geneeskunde VIB-UGent Center for Medical Biotechnology



Assistants



dr. Robbin Bouwmeester



dr. Ralf Gabriels



Toon Callens



Tine Clays



Alireza Nameni



Arthur Declercq



5 credits

contents:

linear regression, logistic regression support vector machine, random forest, gradient boosting, ensemble learning deep neural networks, Al programming with scikit-learn build and evaluate your own models

final competences:

The student is capable of interacting at high level with data analysis specialists.

The student is capable of understanding the specific literature on machine learning based data analysis.



5 credits

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linear regression, logistic regression
support vector machine, random forest, gradient boosting, ensemble learning
deep neural networks, Al
programming with scikit-learn
build and evaluate your own models

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calculation of the examination mark:

(continuous evaluation)
Jupyter notebook project (15%)
microteaching: teach other students about a machine learning method not discussed during the lectures (15%)
2 page report about the obtained Kaggle contest results (20%)

(period-specific evaluation) written exam on 50% of the total score



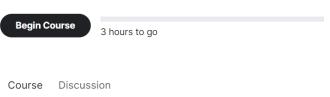
Jupyter notebook project (15%)

Intro to Machine Learning
Intermediate Machine Learning

certificate send to me by email sven.degroeve@ugent.be (subject: kaggle certificate)

Intro to Machine Learning

Learn the core ideas in machine learning, and build your first models.



		Exercise
How Models Work The first step if you're new to machine learning.	PO ₂	
	(8)	
Basic Data Exploration	PS1	$\langle \rangle$
Load and understand your data.		()
Your First Machine Learning Model	PS.	\Diamond
Building your first model. Hurray!	(2)	
Model Validation	PSI	\Diamond
Measure the performance of your model, so you can test and compare alternatives.	(9)	•
Underfitting and Overfitting	₽ 1	$\langle \rangle$
Fine-tune your model for better performance.	(3)	<u> </u>
Random Forests	l [⊗] 1	$\langle \rangle$
Using a more sophisticated machine learning algorithm.	~	
Machine Learning Competitions	PSI	\Diamond
	The first step if you're new to machine learning. Basic Data Exploration Load and understand your data. Your First Machine Learning Model Building your first model. Hurray! Model Validation Measure the performance of your model, so you can test and compare alternatives. Underfitting and Overfitting Fine-tune your model for better performance. Random Forests Using a more sophisticated machine learning algorithm.	The first step if you're new to machine learning. Basic Data Exploration Load and understand your data. Your First Machine Learning Model Building your first model. Hurray! Model Validation Measure the performance of your model, so you can test and compare alternatives. Underfitting and Overfitting Fine-tune your model for better performance. Random Forests Using a more sophisticated machine learning algorithm.



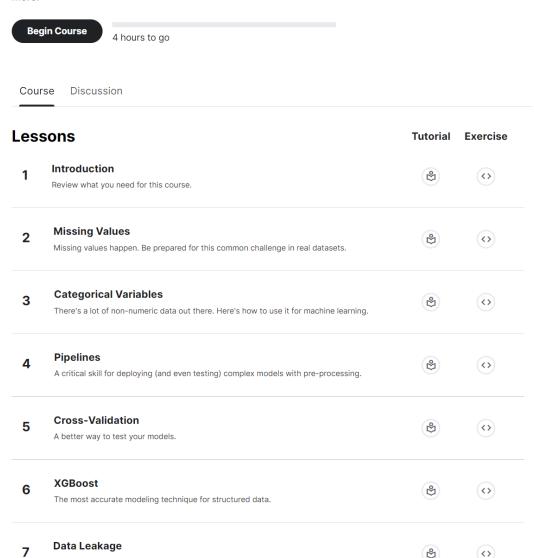
Jupyter notebook project (15%)

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Intermediate Machine Learning

Handle missing values, non-numeric values, data leakage, and more.





Find and fix this problem that ruins your model in subtle ways.

Jupyter notebook project (15%)

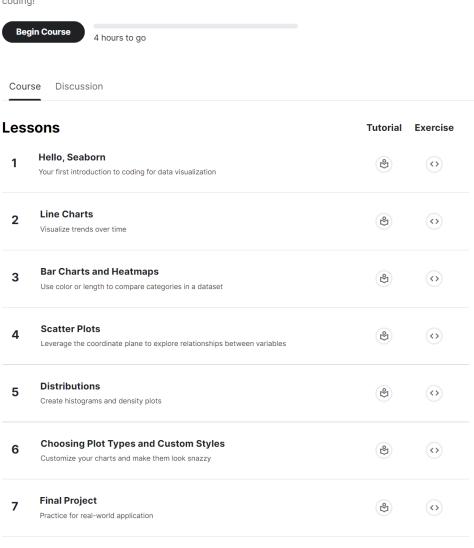
Intro to Machine Learning
Intermediate Machine Learning

certificate send to me by email sven.degroeve@ugent.be (subject: kaggle certificate)

optional:
Data Visualization

Data Visualization

Make great data visualizations. A great way to see the power of coding!



8



Creating Your Own Notebooks

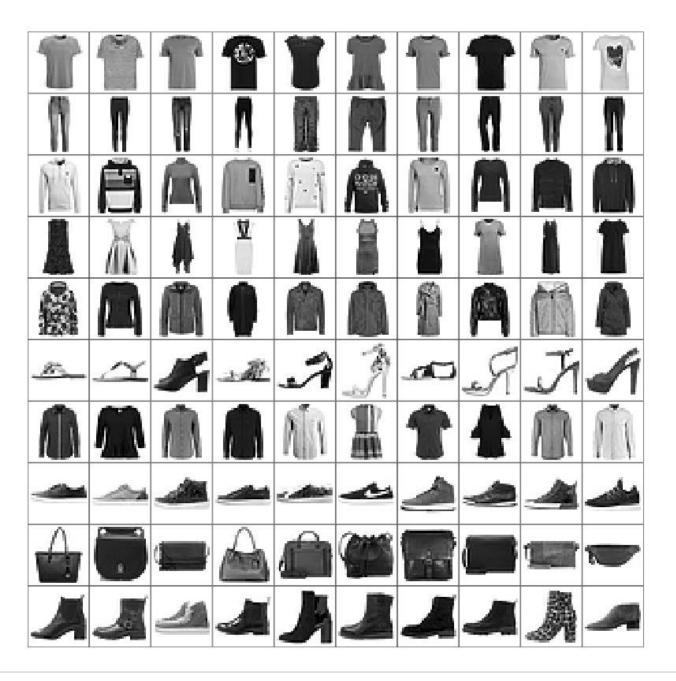
How to put your new skills to use for your next personal or work project

2 page (Calibri (body) font, size 11, A4) report about the obtained Kaggle contest results (20%)





2 page (Calibri (body) font, size 11, A4) report about the obtained Kaggle contest results (20%)

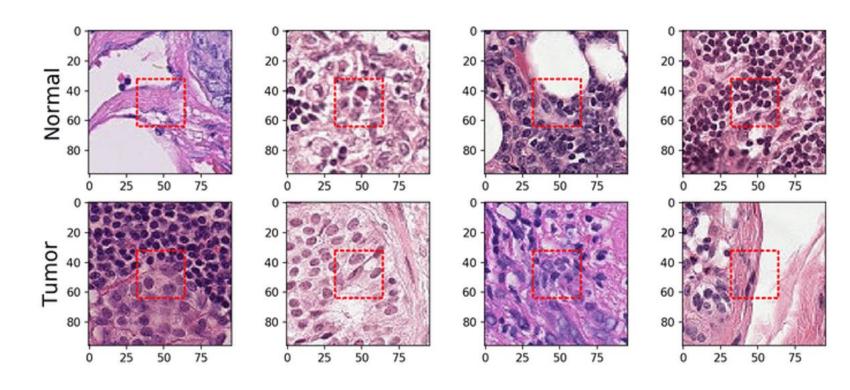




2 page (Calibri (body) font, size 11, A4) report about the obtained Kaggle contest results (20%)

histopathologic scans of lymph node sections

binary label indicating presence of metastatic tissue





2 page (Calibri (body) font, size 11, A4) report about the obtained Kaggle contest results (20%)

format: technical report

should contain:

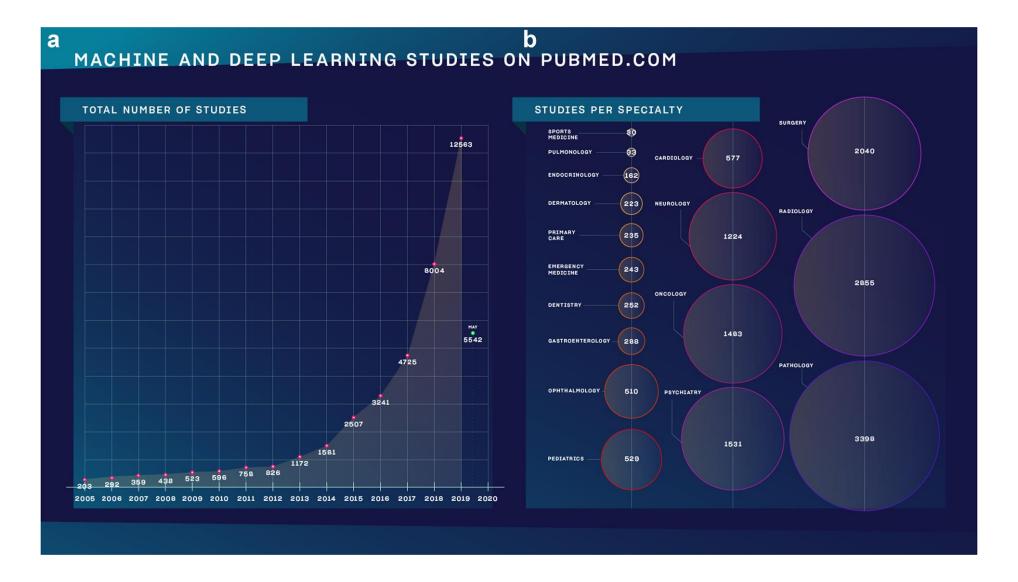
- a clear and concise description of the method(s) used to fit your model(s) (show me you understand what you have done)
- a Table with optimal hyperparameter values
- a Table with evaluation results (both local CV and leaderboard test set)
- a section about what feature(s) are considered most important
- a short Discussion section
- clear plots are of course allowed as well!



microteaching: teach other students about a machine learning method not discussed during the lectures (15%)

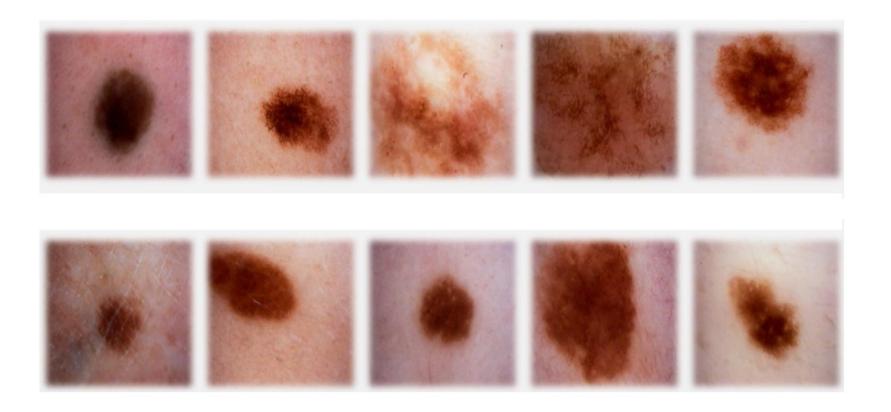


Why follow this course?





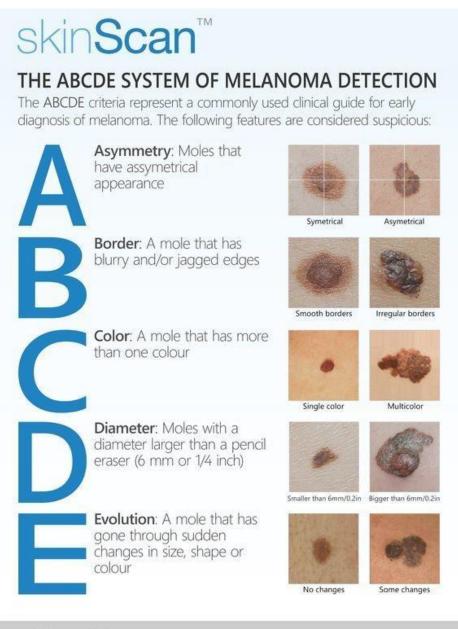
classification



sign of cancer
top row malignant
bottom row benign



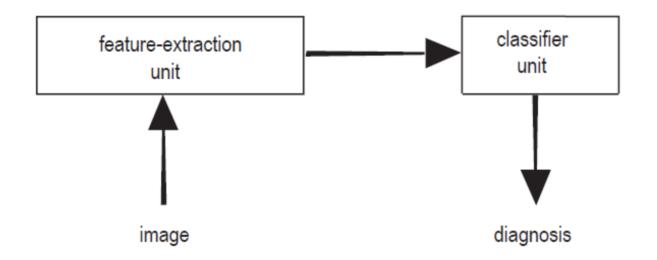
classification



TeleSkin © 2013



classification: terminology

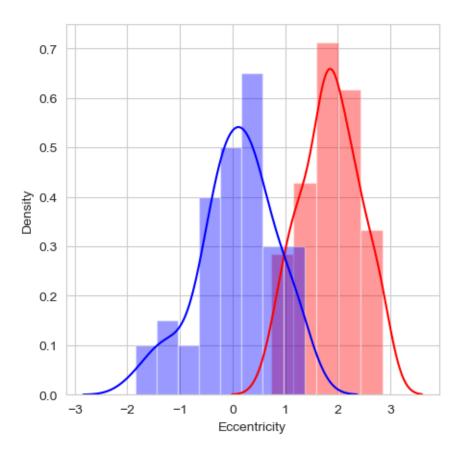


feature extraction: features (a.k.a. properties or attributes)

data set, sample (a.k.a. example, instance or data point), label (a.k.a. target)



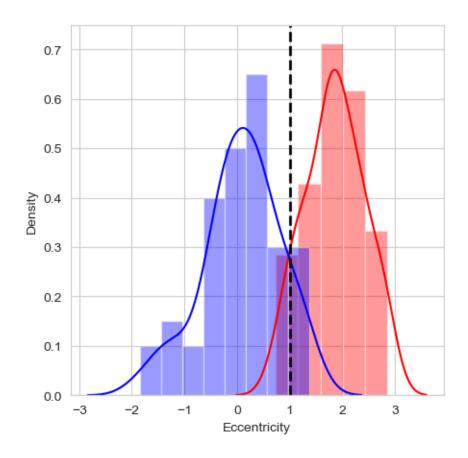
classification: a feature



feature: eccentricity of lesion (how nearly circular the lesion is)



classification: the model

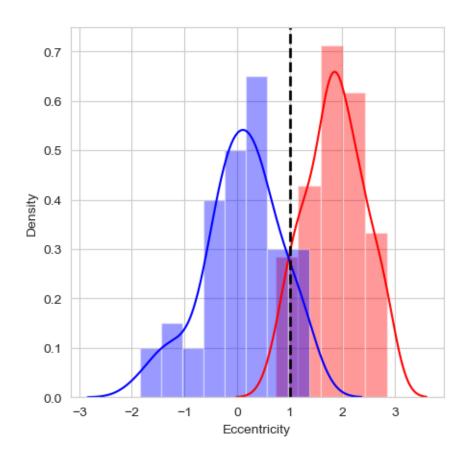


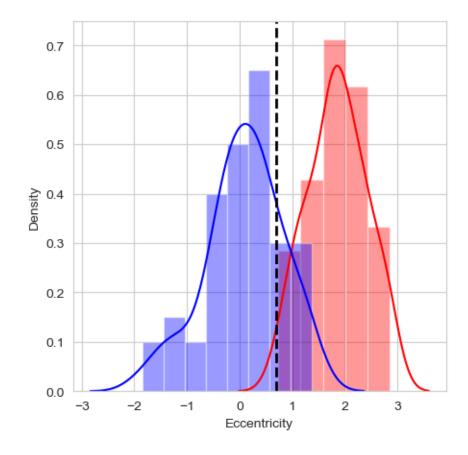
feature: eccentricity of lesion (how nearly circular the lesion is)

model: threshold *t*



classification: the model





feature: eccentricity of lesion (how nearly circular the lesion is)

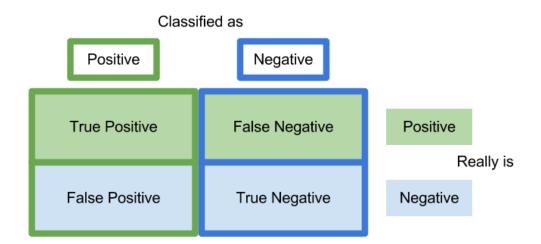
model: threshold t: consequence of the predictions

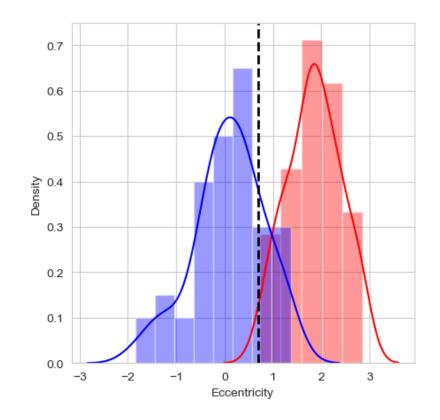


classification: prediction errors

malignant: **positive** class benign: **negative** class

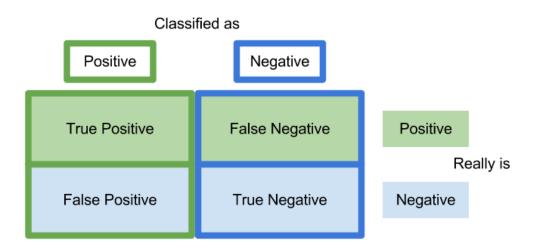
count the number of malignant images with eccentricity value $\geq t$: **true positive** predictions (TP) count the number of malignant images with eccentricity value < t: **false negative** predictions (FN) count the number of benign images with eccentricity value $\geq t$: **false positive** predictions (FP) count the number of benign images with eccentricity value < t: **true negative** predictions (TN)







classification: prediction errors

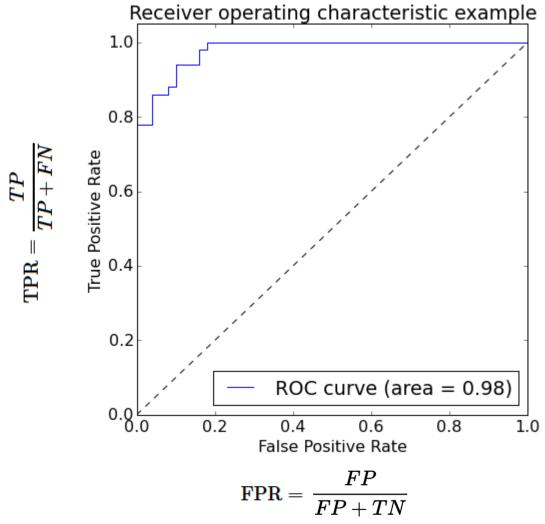


$$\text{accuracy} = \frac{TP + TN}{TP + FP + TN + FN}$$

$$ext{TPR} = rac{TP}{TP + FN}$$

$$ext{FPR} = rac{FP}{FP + TN}$$

classification: prediction errors



model that classifies all images as malignant:

TPR=1 and FPR=1

model that classifies all images a benign:

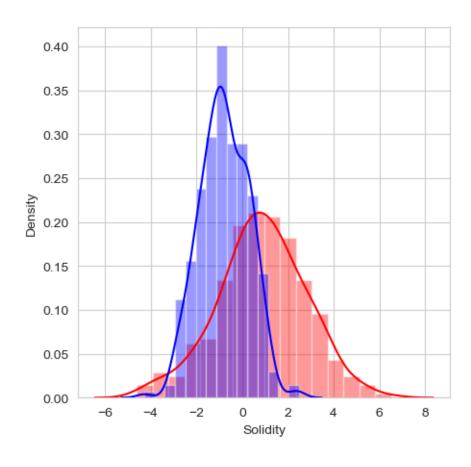
TPR=0 and FPR=0

vary threshold *t*

Area Under the Curve (AUC)



classification: multi-dimensional



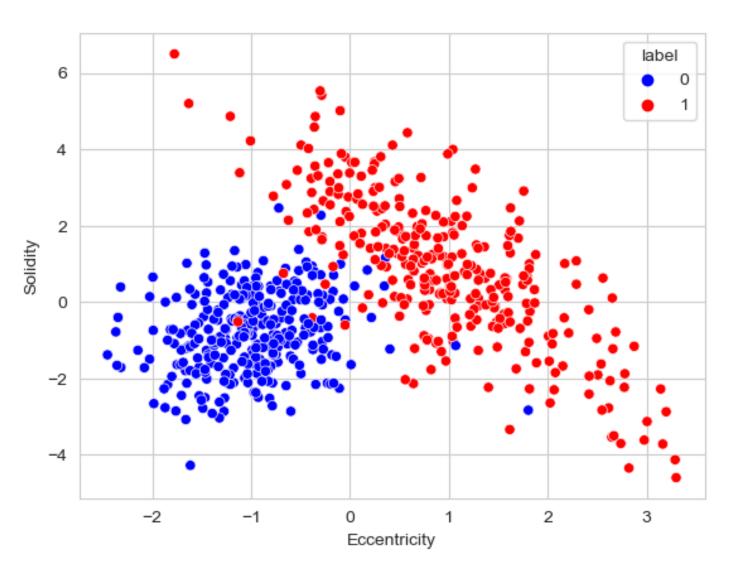
add another feature?

feature vector X

Euclidean vector space



classification: multi-dimensional

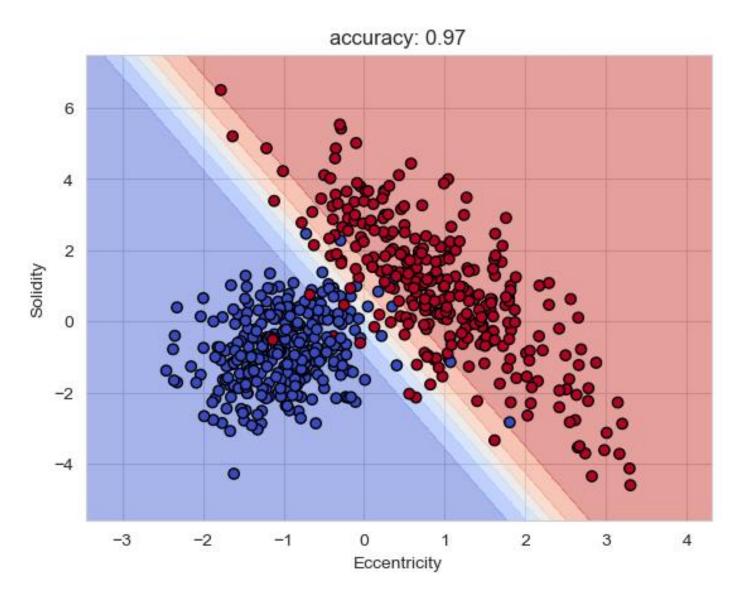


feature vector X

Euclidean vector space



classification: multi-dimensional



linear decision boundary

blue region malignant, red region benign

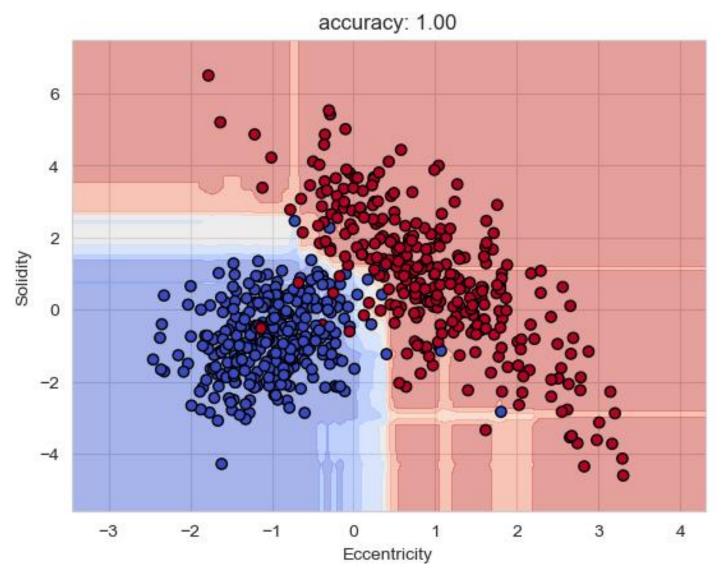
yet more features

can't look at the decision boundary

more complex



classification: model complexity



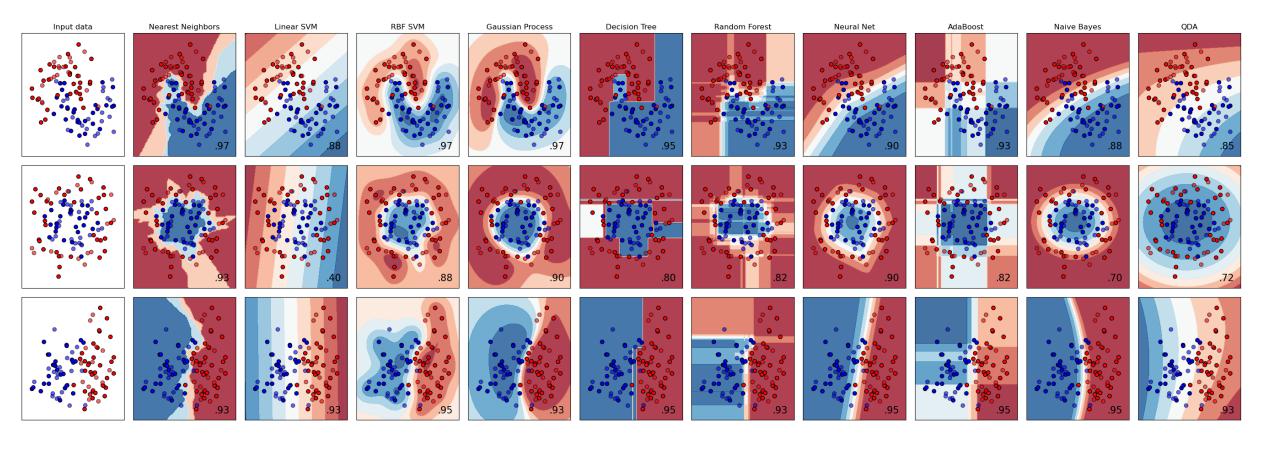
unseen external images

generalization

overfitting



scikit-learn





data normalization

make all features same scale

Eccentricity [0,100], Solidity [-5,7]

weights all features equally in their representation

standardization

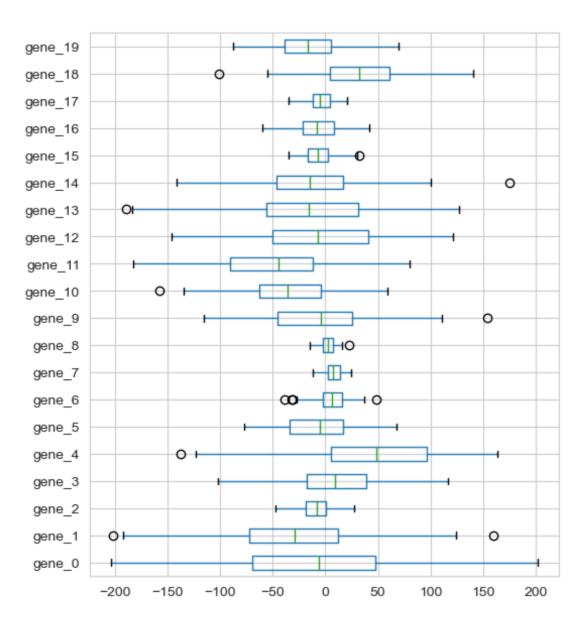
$$\mu = 0$$
 $\sigma = 1$

min-max scaling: scale the features to a fixed range

$$x_{norm} = rac{x - \mu}{\sigma}$$

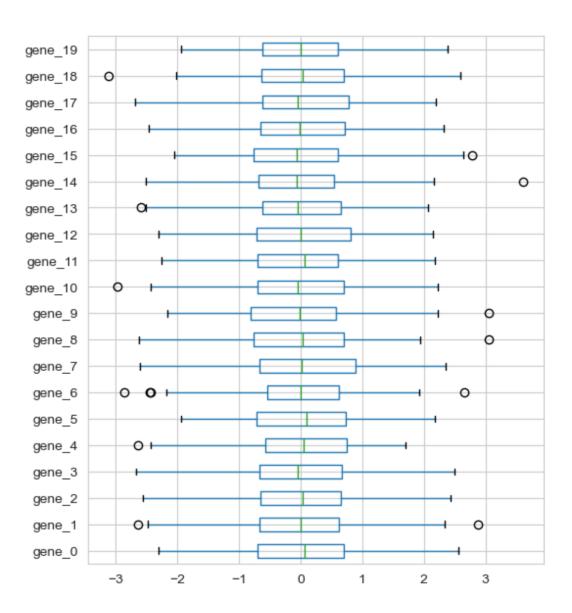
$$x_{norm} = rac{x - x_{min}}{x_{max} - x_{min}}$$

data normalization





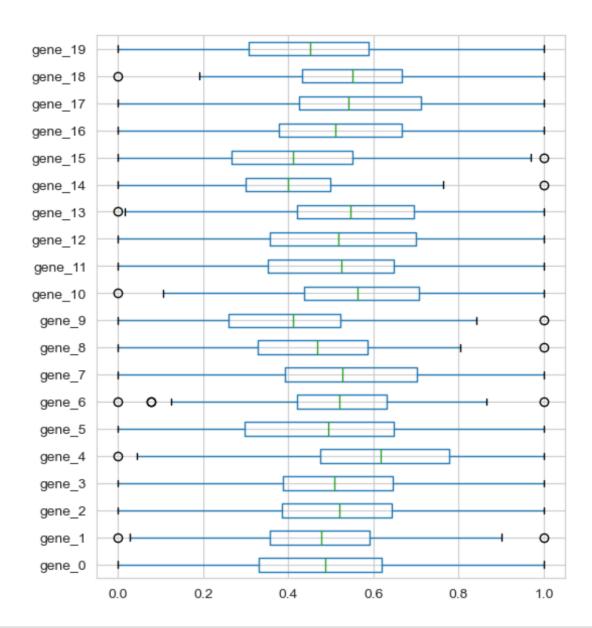
data normalization: standardization



$$x_{norm} = rac{x-\mu}{\sigma}$$



data normalization: min-max scaling



$$x_{norm} = rac{x - x_{min}}{x_{max} - x_{min}}$$

