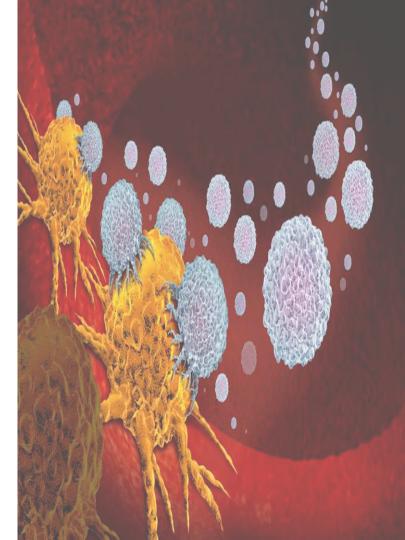


- Cancer mortality can be reduced if cases are detected and treated early
- *Is it possible to predict it?*
- Which factors causes cervical cancer?

• Hospital agency meeting



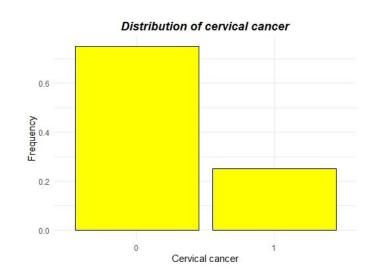
1. Data cleaning

2. Exploratory Descriptive Analysis

- 3. Variables importance
- 4. Parametric method

dataset

- Data collection from <u>surveys</u>: demographic information, habits and historical medical records
- Cervical Cancer target variable: 25% of the patients diagnosticated with cervical cancer: Healthy patients 660 out of 880
- *Uci Machine Learning Repository*: 880 observations, 25 variables



Predictors

Q U A 1. Age N 2. Number of sexual partners 3. Years old when first having sex T 4. Years of smoke 5. Years of hormonal contraceptive Т 6. Years of intra-uterine disease A 7. Number of sexually trasmitted diseases T 8. Number of diagnosis 9. Number of pregnancies V \mathbf{E}

1. Smoke (0-1) 2. Hormonal contraceptive (0-1) 3. Intra-uterine dispositive (0-1) 4. Sexually trasmitted disease (0-1) 5. Condylomatosis (0-1) 6. Syphilis (0-1) 7. Genital herpes (0-1) 8. *Molluscum contagious* (0-1) 9. HIV (0-1) 10. Hepatitis B (0-1) 11.HPV (0-1) 12. DX test (0-1) 13. Cancer (0-1) 14. Schiller test (0-1) 15. Cervical cytology test (0-1) 16. Hinselmann test (0-1)

Data cleaning (1)

Possible distortive variables:

- **Hinselmann test:** to detect **cervical** tumors when they were relatively small.
- Schiller test: iodine solution is applied to the cervix in order to diagnose cervical cancer.
- Cervical cytology: to detect abnormal or potentially abnormal cells from the uterine cervix

Data cleaning (2)

Missing values:

Replace with median:

Number of sexual partner: 28

Years of first sex: 7

Number of pregnancies: 56

Years of smoke: 13

Years of hormonal contraceptive: 108

Years of IUD: 117

Replace with mode:

Smoke: 13

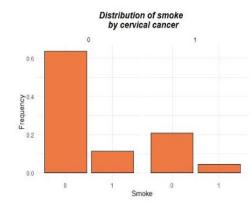
Hormonal contraceptive: 108

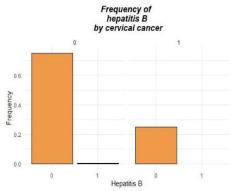
IUD: 117 HIV: 105

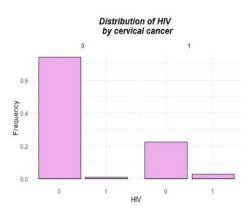
- 1. Data cleaning
- 2. Exploratory Descriptive Analysis

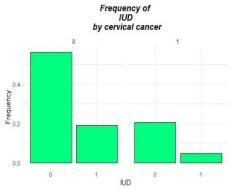
- 3. Variables importance
- 4. Parametric method

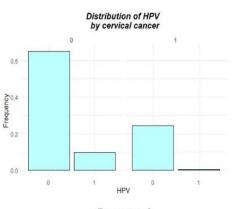
EDA: Qualitative predictors

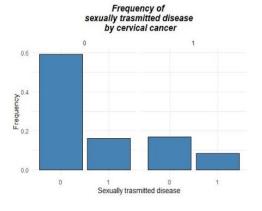




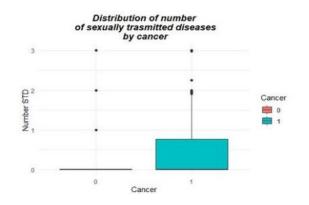


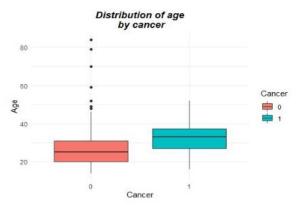


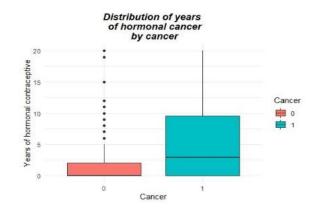


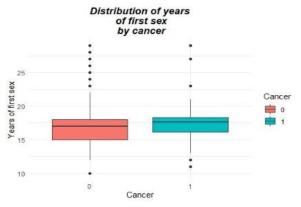


EDA: Quantitative predictors



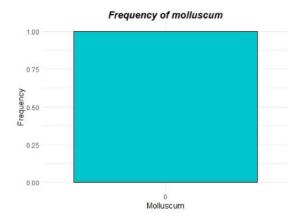


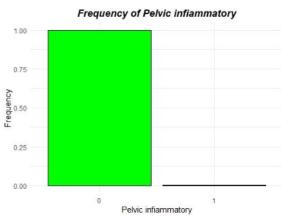


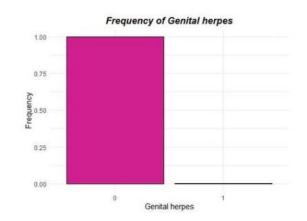


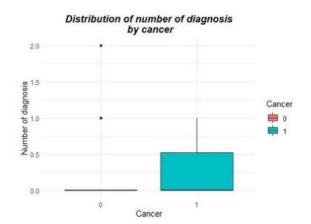
EDA: Not explicative variables

We decide to avoid them:



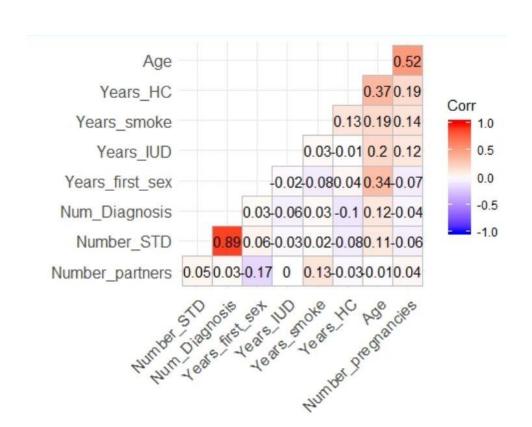






EDA: Correlations

- ❖ Not relevant correlations
- Number of sexually trasmitted diseases correlated with number of cancer diagnosis
- ❖ *Age with number of pregnancies*



- 1. Data cleaning
- 2. Exploratory Descriptive Analysis
- 3. Variables importance
- 4. Parametric method

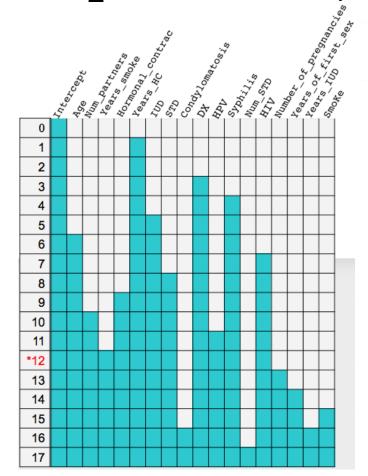
Exploring variables importance(1)

Variables selection:

- ☐ Try with different method: forward, backward, bothwise
- ☐ *Best is the forward with AIC: 727.11*

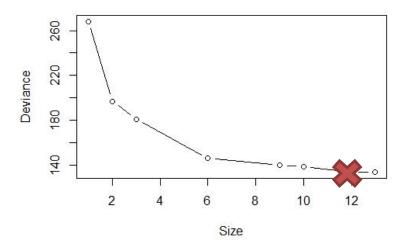
Coefficients:

	Estimate	Std. Error	z value	Pr(> z)	
(Intercept)	-2.41645	0.40941	-5.902	3.59e-09	* * *
Years HC	0.19522	0.02460	7.936	2.09e-15	* * *
Number_STD	0.64331	0.25285	2.544	0.01095	*
DX1	2.64243	0.38847	6.802	1.03e-11	* * *
Syphilis1	-16.61544	496.10492	-0.033	0.97328	
IUD1	-0.84347	0.30553	-2.761	0.00577	* *
Age	0.02995	0.01205	2.486	0.01291	*
HIV1	0.60233	0.51202	1.176	0.23944	
STD1	1.24189	0.45106	2.753	0.00590	* *
Hormonal_contrac1	-0.43646	0.24035	-1.816	0.06938	
Number_partners	-0.14777	0.07108	-2.079	0.03762	*
HPV1	-1.36277	0.70356	-1.937	0.05275	
Years smoke	0.04216	0.02172	1.941	0.05230	



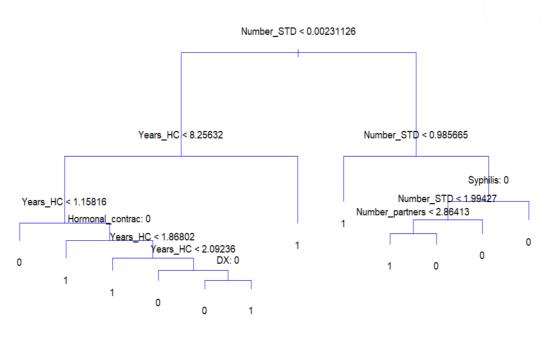
Exploring variables importance (2)

Tree:



Cross Validation to select the best split: 12

- Number of sexually trasmitted diseases;
- Years of hormonal contraceptive;
- Number of partners;
- Syphilis
- DX

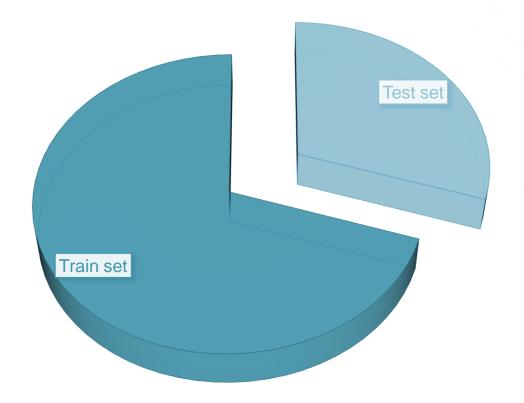


Misclassification error rate 10.68%

- 1. Data cleaning
- 2. Exploratory Descriptive Analysis
- 3. Variables importance
- 4. Parametric method

TRAIN AND TEST SET

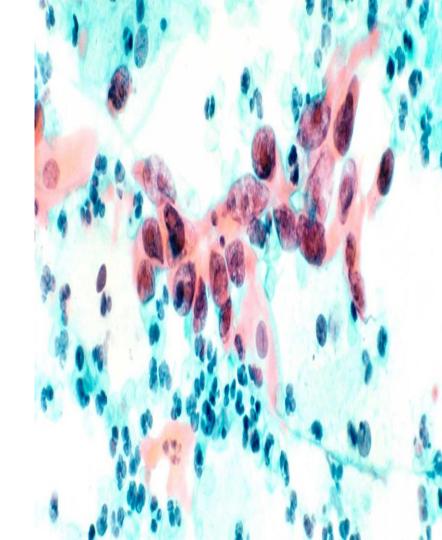
- 70% training set
- 30% test set



GLM (1)

Based on forward selection:

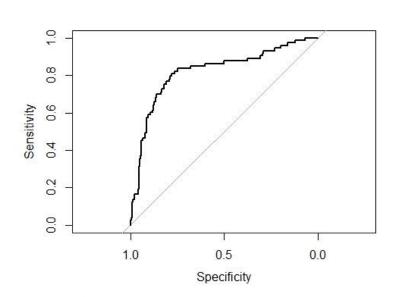
	Exp (coeff.)	Significativity
Intercept	0.156705	***
Years HC	1.25446	***
Number STD	2.240323	*
DX1	18.88446	***
Syphilis1	5.623143 e-08	
IUD1	0.6178992	
Age	1.016901	
HIVI	3.494883	
STD1	3.386815	*
НС	0.4610955	*
Number_partners	0.8085199	*
HPV1	0.4661583	
Years_smoke	1.0365	



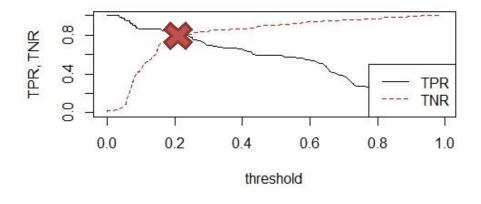
GLM(2)

Choice of the threshold:

AUC: 82.53%



THRESHOLD: 0.1832281



	False	True
0	172	49
1	13	60

- Specificity: 77.40%
- Sensitivity: 83.56%
- Accuracy: 79%

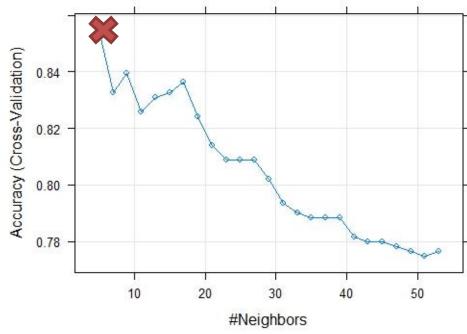
- 1. Data cleaning
- 2. Exploratory Descriptive Analysis

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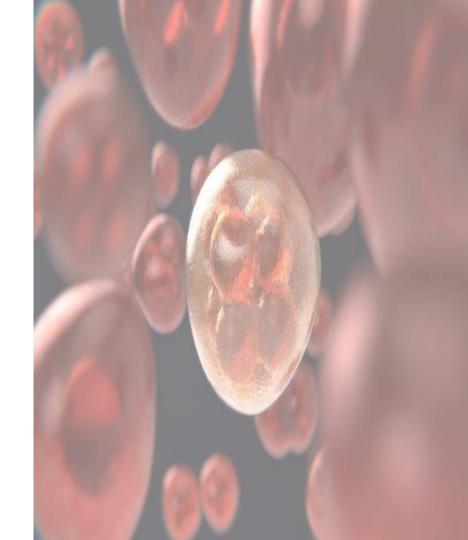
KNN

10-fold cv for the best K:

K=5



Accuracy 85%



RANDOM FOREST (1)

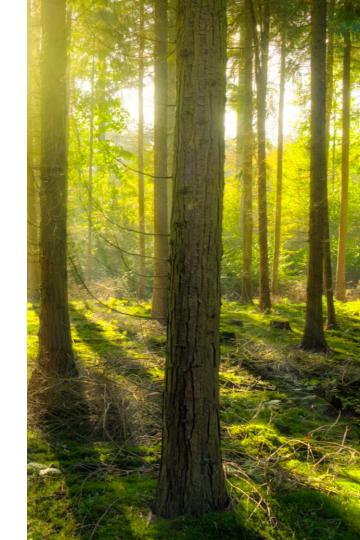
 Aggregating many decision trees improves the predictive performance

Bagging reduce variance in trees:
 m=p

• Decorrelating the trees

<u>considering only a subset m of the</u>

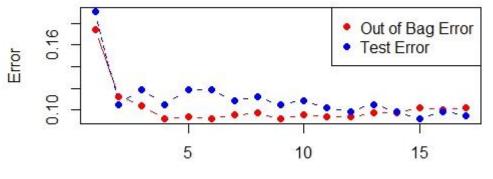
<u>predictors</u> m<p



RANDOM FOREST (2)

How to select the best number of predictors at each split?

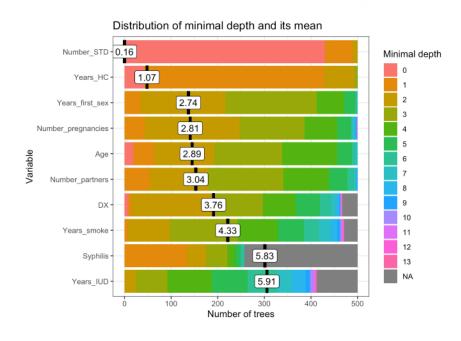
Try with all possible values of "mTry":



Number of Predictors Considered at each Split

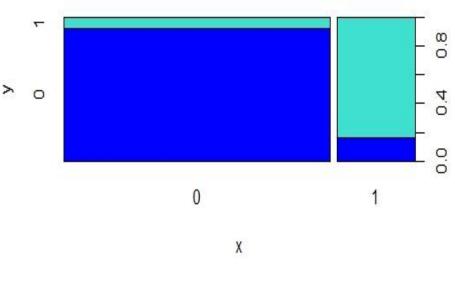
The best is m=15

- Out of bag error (OOB): 9.22%
- *Test error:* 9.18%

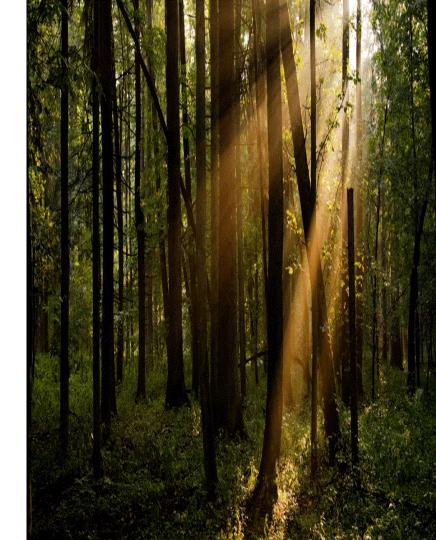


RANDOM FOREST (3)

Predictions m=15



Accuracy: 91%



CONCLUSION AND IMPROVEMENTS

- 1. Cancer most influencing variables: number of sexually transmitted diseases, years of hormonal contraceptive
- 2. Best prediction with random forest

GLM	KNN	RANDOM FOREST
79 %	85 %	91 %

- 3. Try different methods for data cleaning
- 4. Cross validation for random forest and GLM threshold: to improve '1' prediction
- 5. Try Neural Network

REFERENCES

- Gareth James, Daniela Witten, Trevor Hastie, Robert Tibshirani,
 "An Introduction to Statistical Learning", Springer Science+Business Media New York
 2013
- Muhammed Fahri Unlersen1, Kadir Sabanci2, Muciz Özcan1, "Determining Cervical Cancer Possibility by Using Machine Learning Methods",
 International Journal of Latest Research in Engineering and Technology, December 2017