

# Mini-project

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# Agenda

- 1 Process of choosing datasets
- 2 Selected datasets
- 3 Preprocessing of datasets
- 4 Models
- 5 Training
- 6 Results
- 7 Comparison/Discussion
- 8 Conclusion

# Datasets and preprocessing

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Pipeline for selecting and processing the data

# Dataset selection process

Criteria - easy to deal with:

- Based on article
- No special preprocessing
- Limited size

Result: 22 selected datasets

Algerian forest fire:

Classes	
fire	131
not fire	101
fire	4
fire	2
not fire	2
not fire	1
not fire	1
not fire	1

Example of not included dataset.

# Selected datasets

Dataset	No. samples	No. features	Missing values	Majority class %	No. classes	UCI id
acute_inflamations	120	7	No	58.3	2	184
balance_scale	625	4	No	46.1	3	12
balloons	<b>16</b>	4	No	56.3	2	13
breast_cancer_wisconsin_diagnostic	569	30	No	62.7	2	17
car_evaluation	1728	6	No	70.0	4	19
congress_voting_records	232	16	<b>Yes</b>	61.4	2	105
credit_approval	653	15	<b>Yes</b>	55.5	2	27
ecoli	336	7	No	42.6	<b>8</b>	39

# Selected datasets

Dataset	No. samples	No. features	Missing values	Majority class %	No. classes	UCI id
fertility	100	9	No	<b>88</b>	2	244
habermans_survival	306	<b>3</b>	No	73.5	2	43
hayes_roth	132	4	<b>Yes</b>	31.9	4	44
heart_disease	297	13	<b>Yes</b>	54.1	5	45
ilpd	579	10	<b>Yes</b>	71.4	2	225
iris	150	4	No	33.3	3	53
lenses	24	<b>3</b>	No	62.5	3	58
mammographic_mass	830	5	<b>Yes</b>	53.7	2	161

# Selected datasets

Dataset	No. samples	No. features	Missing values	Majority class %	No. classes	UCI id
mushroom	5644	22	Yes	51.8	2	73
spect_heart	267	22	No	79.4	2	95
spectf_heart	267	44	No	79.4	2	96
statlog	1000	20	No	70	2	144
wine_quality	6497	11	No	43.7	7	186
zoo	101	16	No	40.6	7	111

# Selected datasets

Dataset	No. samples	No. features	Missing values	UCI id	Our id
acute_inflamations	120	7	No	184	1
balance_scale	625	4	No	12	2
balloons	<b>16</b>	4	No	13	3
breast_cancer_wisconsin_diagnostic	569	30	No	17	4
car_evaluation	1728	6	No	19	5
congress_voting_records	232	16	<b>Yes</b>	105	6
credit_approval	653	15	<b>Yes</b>	27	7
ecoli	336	7	No	39	8



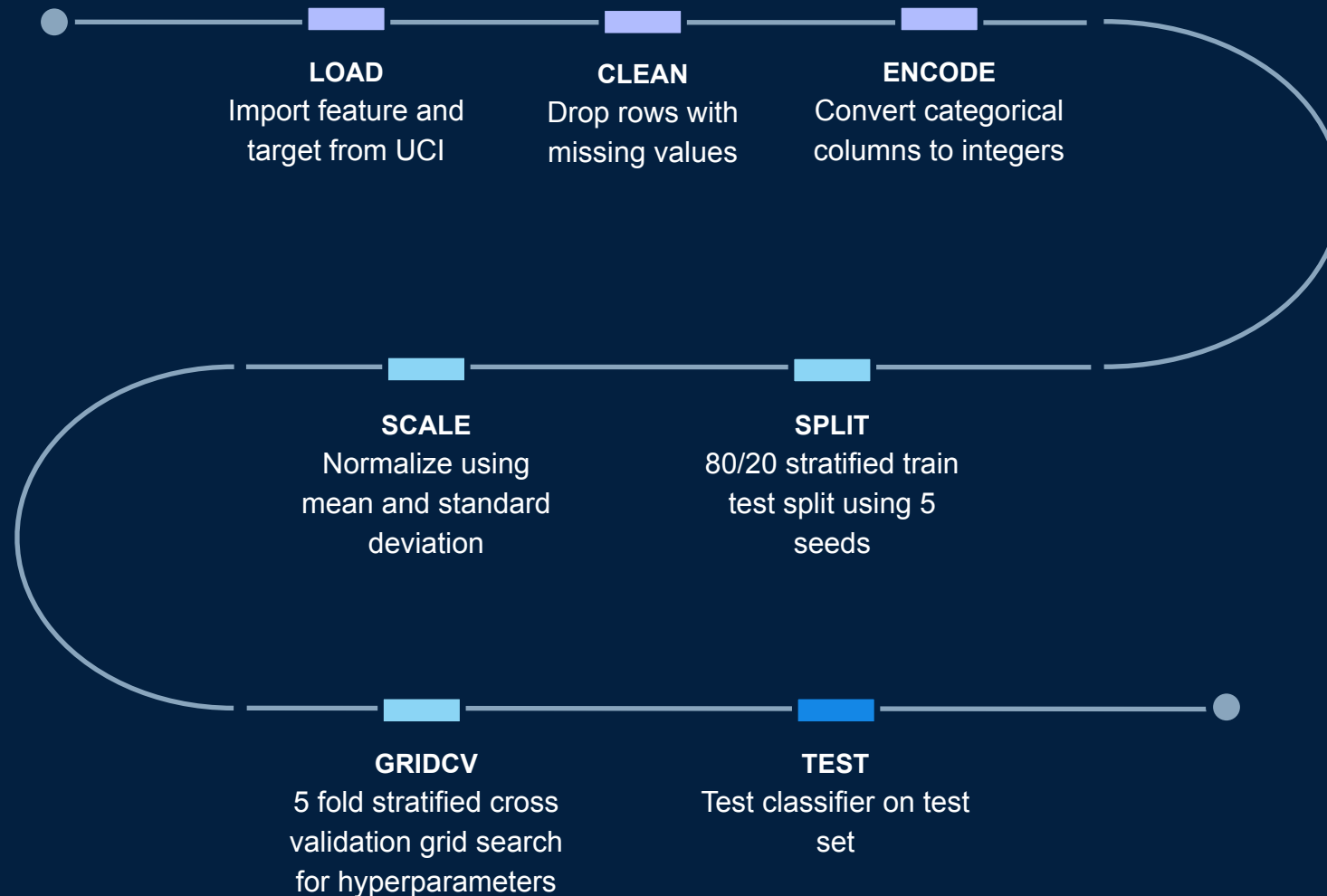
# Selected datasets

Dataset	No. samples	No. features	Missing values	UCI id	Our id
fertility	100	9	No	244	9
habermans_survival	306	3	No	43	10
hayes_roth	132	4	Yes	44	11
heart_disease	297	13	Yes	45	12
ilpd	579	10	Yes	225	13
iris	150	4	No	53	14
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# Selected datasets

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spectf_heart	267	<b>44</b>	No	96	19
statlog	1000	20	No	144	20
wine_quality	<b>6497</b>	11	No	186	21
zoo	101	16	No	111	22

# Overview: pipeline



# Preprocessing



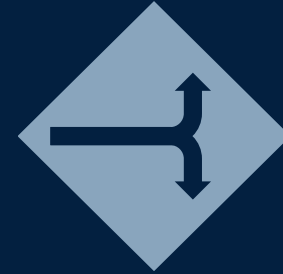
## Cleaning

Remove rows with missing values.



## Encoding

Convert categorical features into numerical.



## Splitting

Stratified 80-20 train-test split.



## Scaling

Normalize using mean and standard deviation.

Based on article

# Models and Training

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For supervised and unsupervised classification

## Models: supervised

1

Random forest

2

Support vector  
machine

3

Logistic  
regression

4

K-nearest  
neighbors

5

Gaussian naive  
bayes

6



# Hyperparameter optimization

## Random Forest

```
random_forest_params = [  
    {"random_forest__n_estimators": [100, 500],  
     "random_forest__max_depth" : [5, 10, 15]  
    }  
]
```

## Logistic Regression

```
log_reg_params = [  
    {'log_reg__solver': ["lbfgs", "saga"],  
     'log_reg__penalty': ['l2'],  
     'log_reg__C' : np.logspace(-3,3,7),  
     'log_reg__max_iter' : [100,1000,2500]  
    }  
]
```

## SVM

```
svm_params = [{"svm__C": [0.1, 1, 10, 100, 1000],  
               'svm__gamma': [1, 0.1, 0.01, 0.001, 0.0001],  
               'svm__kernel': ['rbf']} ]
```

## KNN

```
knn_params = [{"knn__n_neighbors": [3, 5, 7, 9],  
               'knn__weights': ['uniform', 'distance'],  
               'knn__leaf_size': [15, 20]}]
```

## Gaussian naive bayes

```
gnb_params = [  
    {'gnb__var_smoothing': np.logspace(0,-9, num=10)  
    }  
]
```

# Models: unsupervised

1

Agglomerative  
clustering

2

Affinity  
propagation

3

K-means

4





# Hyperparameter optimization

## Affinity Propagation

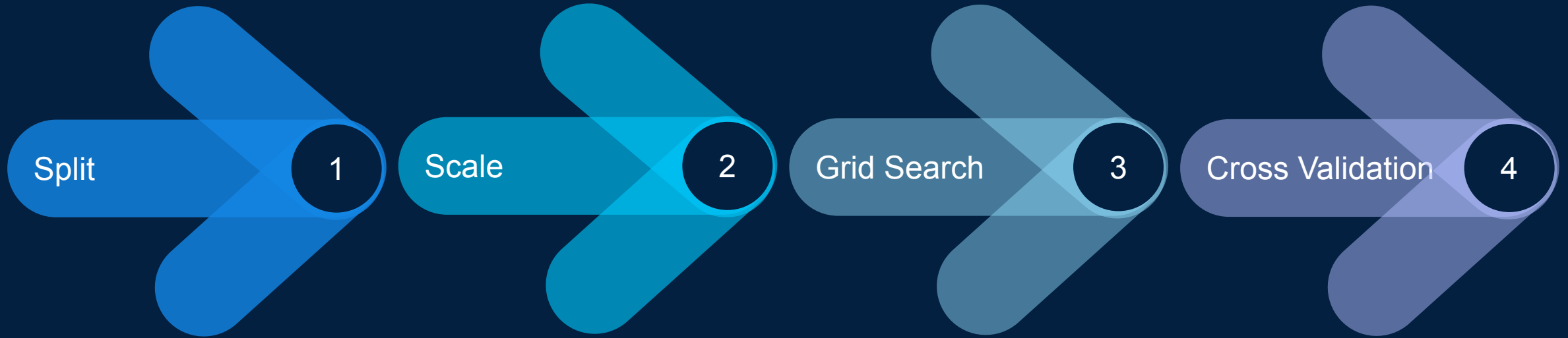
```
affinity_propagation_params = [  
    {"damping": [0.5, 0.7]},  
]
```

## Agglomerative Clustering

```
metrics = ["euclidean", "l1", "l2", "manhattan"]  
linkages = ["complete", "average", "single"]  
pca_options = [True, False]
```

```
score = 0.8 * val_acc + 0.2 * train_acc  
if score > best_score:  
    best_params = params  
    best_score = score
```

# Training



For K-means and affinity propagation: mode mapping

For agglomerative clustering: Hungarian algorithm mapping, no single prediction, no cross validation (80/20 train/validation split)

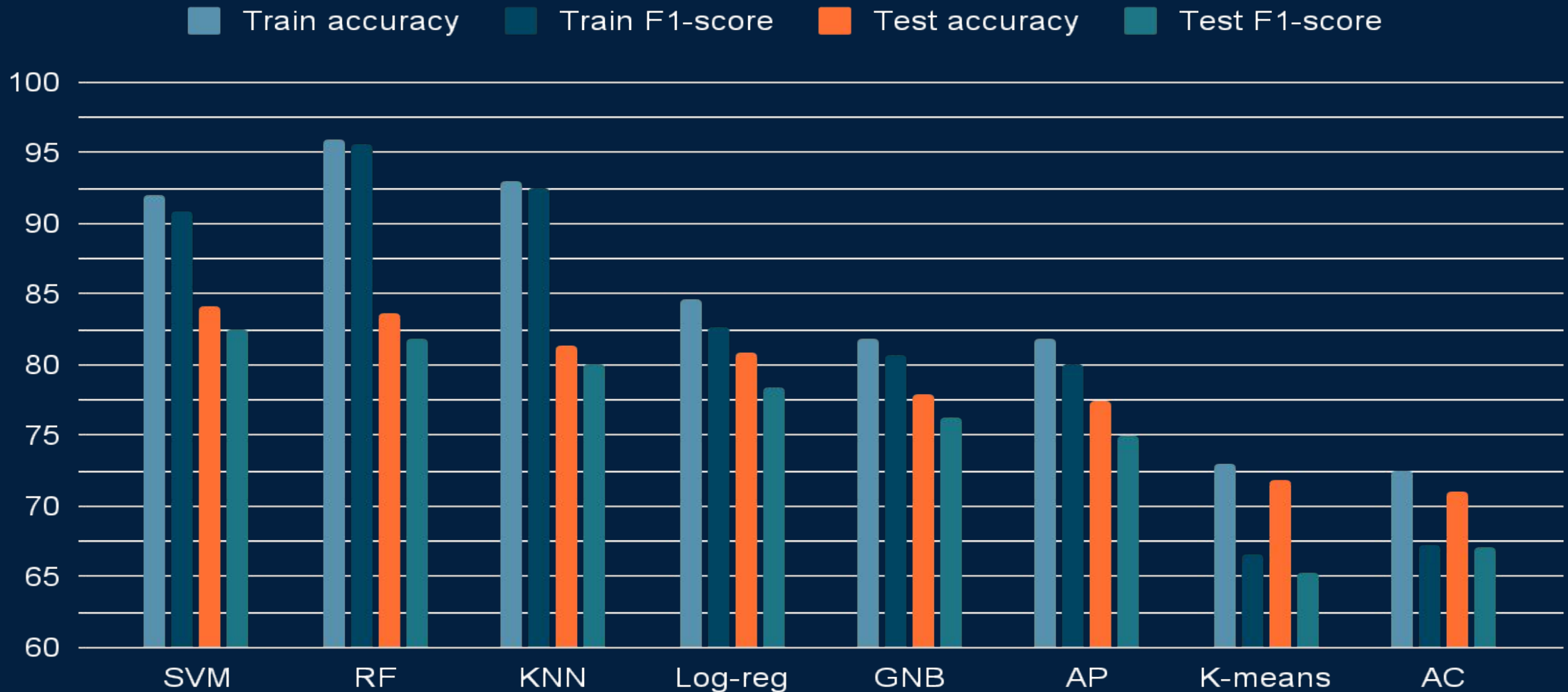
# Results and Discussion

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Our results in comparison to the article

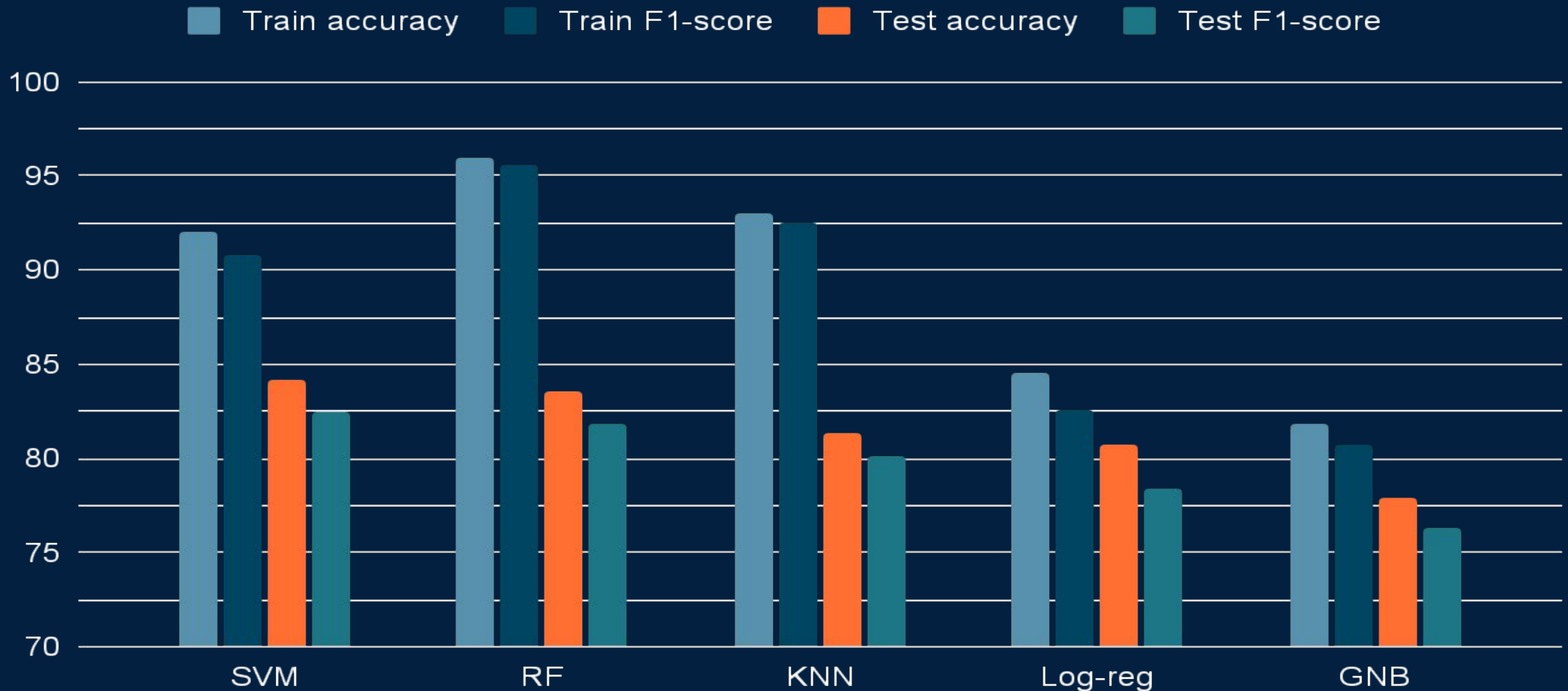
# Results

## Average metrics



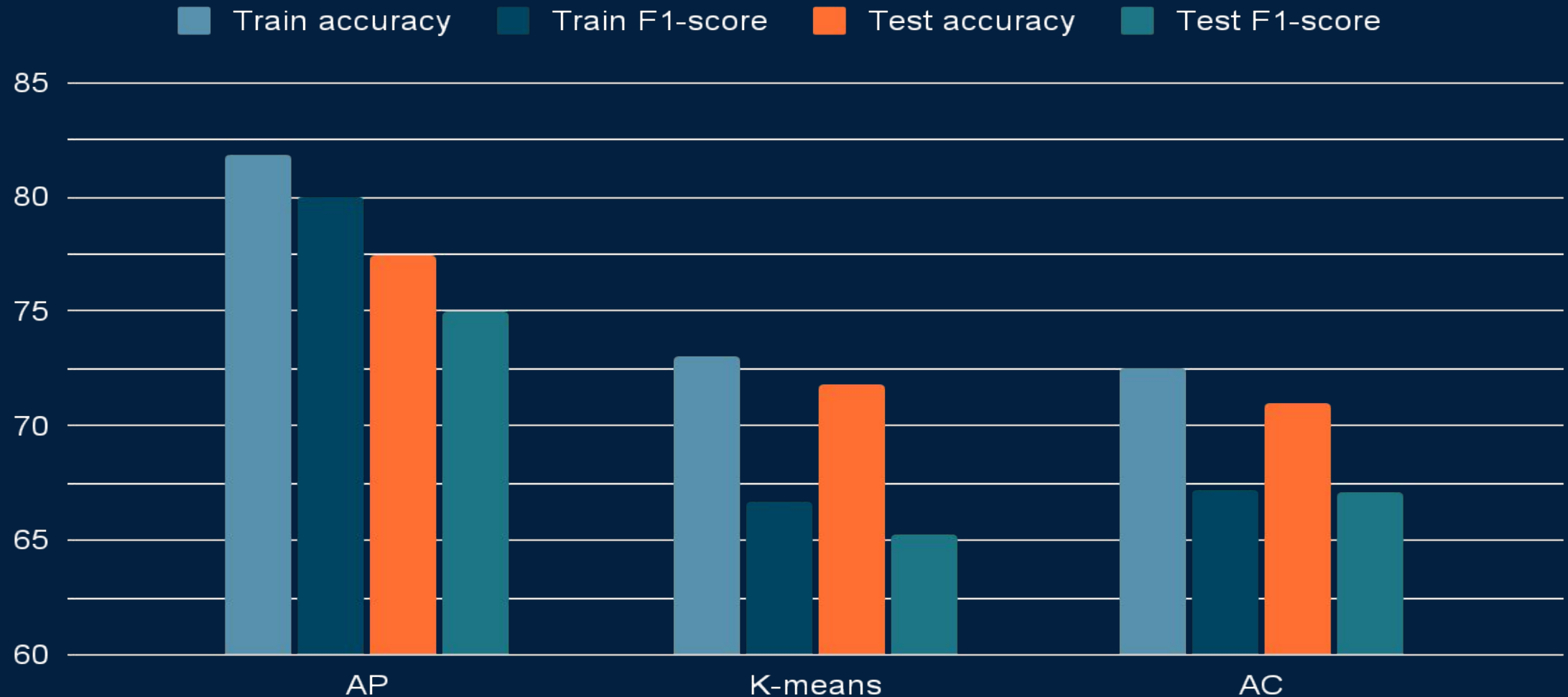
# Results: supervised

## Average metrics supervised



# Results: unsupervised

## Average metrics unsupervised



# Results

To see all results:

We saved all results in a tensorboard logger

Our runs are saved in our [GitHub](#)

Command to start (given tensorboard installed, replace runs/ with path to folder, might have to try different port number)

```
tensorboard --logdir runs/ --port 6006 --samples_per_plugin images=22
```

Regex for colours:

```
(knn|kmeans|svm|random|gnb|log_reg|agglo|affini)
```

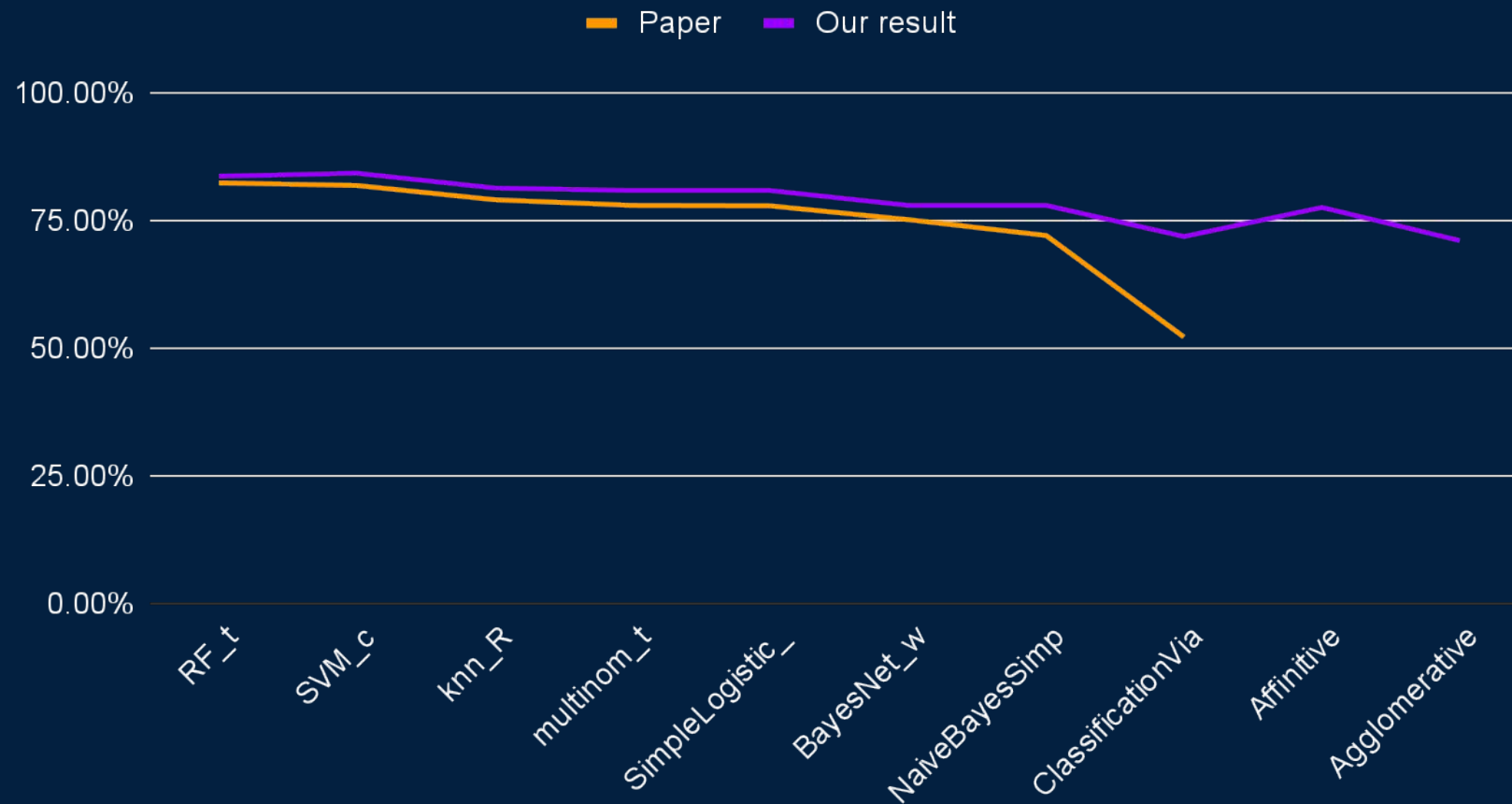
# Comparison/discussion

	Classifier name	Paper (average accuracy)	Our result
Random Forest	RF_t	82.3 %	83.6 %
Support Vector Machine	SVM_c	81.8 %	84.2 %
K-Nearest Neighbor	knn_R	79.0 %	81.3 %
Logistic Regression	multinom_t	77.9 %	80.8 %
	SimpleLogistic_w	77.8 %	
Naive Bayes	BayesNet_w	75.1 %	77.9 %
	NaiveBayesSimple_w	72.0 %	
Clustering (K-Means)	ClassificationViaClustering_w	52.1%	71.8 %
Affinity Propagation	/	/	77.5 %
Agglomerative Clustering	/	/	71



# Comparison/discussion

Paper vs our results (Test accuracy per classifier)



# Conclusion

- implemented 8 classifiers for 21 datasets, common preprocessing and CV
- reach similar but slightly better results than the paper
- Difficulties with imbalanced datasets => stratified sampling, F1-score
- Supervised classifiers were easier to implement with the pipeline that is provided by sklearn
- learned about different techniques how to remap labels for unsupervised classifiers
- Future additions:
  - removing highly correlated columns
  - adding more classifiers
  - adding more possible combinations of hyperparameters
  - adding more complex datasets

Thank you for your attention!

# Kontakt

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Github: [https://github.com/Dropptimus/D7041E\\_Mini\\_project](https://github.com/Dropptimus/D7041E_Mini_project)  
=> full\_pipeline\_fixed\_metrics\_multiple\_seed.ipynb

