

ROBERT v 1.0.5 2024/09/06 19:23:24

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ROBERT SCORE

This score is designed to analyze the predictive ability of the models using different metrics.

No PFI (all descriptors):

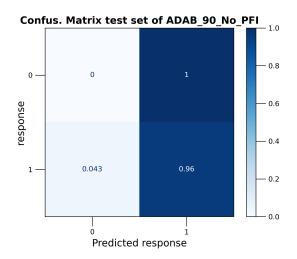
ML model: ADAB

Proportion Train: Validation: Test = 81:9:10

VERY WEAK

The model has a score of 2/10

- The test set shows an accuracy of 0.79
- The test set uses 259:115 points:descriptors
- The test set passes 1 VERIFY tests
- The test set passes 0 y-mean/y-shuffle



Train : Accuracy = 0.83, F1 score = 0.9, MCC = 0.46 Valid.: Accuracy = 0.96, F1 score = 0.98, MCC = 0.8 Test: Accuracy = 0.79, F1 score = 0.88, MCC = -0.09

PFI (only important descriptors):

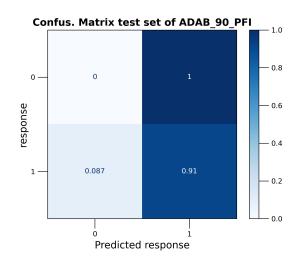
ML model: ADAB

Proportion Train: Validation: Test = 81:9:10



The model has a score of 4/10

- The test set shows an accuracy of 0.75
- The test set uses 259:3 points:descriptors ••
- The test set passes 1 VERIFY tests
- The test set passes 0 y-mean/y-shuffle



Train: Accuracy = 0.79, F1 score = 0.88, MCC = 0.28 Valid.: Accuracy = 0.92, F1 score = 0.96, MCC = 0.0 Test: Accuracy = 0.75, F1 score = 0.86, MCC = -0.13

Score thresholds (detailed in https://robert.readthedocs.io/en/latest/Score/score.html)

Accuracy

● Accuracy > 0.85

- 0.85 > Accur. > 0.70
- Accur. < 0.70

Outliers

- Excluded in classif.

Points:descriptors

- > 10:1 p:d ratio
- 10:1 > p:d ratio > 3:1
- p:d ratio < 3:1

VERIFY tests

- • y-shuffle & y-mean
- 5-fold CV & onehot
- (all tests failed)

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Some tips to improve the score

Adding meaningful descriptors or replacing/deleting the least useful descriptors used might help. Feature importances are gathered in the SHAP and PFI sections of the /PREDICT/PREDICT data.dat file.

How to predict new values with these models?

- 1. Create a CSV database with the new points, including the necessary descriptors.
- 2. Place the CSV file in the parent folder (i.e., where the module folders were created)
- 3. Run the PREDICT module as 'python -m robert --predict --csv_test FILENAME.csv'.
- 4. The predictions will be shown at the end of the resulting PDF report and will be stored in the last column of two CSV files called MODEL_SIZE_test(_No)_PFI.csv, which are in the PREDICT folder.



REPRODUCIBILITY

This section provides all the instructions to reproduce the results presented.

- 1. Download these files (the authors should have uploaded the files as supporting information!):
 - CSV database (AQME-ROBERT_lipros_data.csv)
- 2. Install and adjust the versions of the following Python modules:
 - Install ROBERT and its dependencies: conda install -c conda-forge robert
 - Adjust ROBERT version: pip install robert==1.0.5
 - scikit-learn-intelex: not installed

(if scikit-learn-intelex is installed, slightly different results might be obtained)

3. Run ROBERT using this command line in the folder with the CSV database:

python -m robert --y "response" --csv_name "AQME-ROBERT_lipros_data.csv" --type "clas" --names "code_name"

4. Execution time, Python version and OS:

Originally run in Python 3.10.14 using Linux #3672-Microsoft Fri Jan 01 08:00:00 PST 2016

Total execution time: 90694.58 seconds (the number of processors should be specified by the user)

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TRANSPARENCY

This section contains important parameters used in scikit-learn models and ROBERT.

1. Parameters of the scikit-learn models (same keywords as used in scikit-learn):

No PFI (all descriptors): PFI (only important descriptors):

sklearn model: AdaBoostClassifier sklearn model: AdaBoostClassifier

random state: 43 random_state: 43 names: code name names: code name n estimators: 20 n estimators: 20 learning rate: 0.5 learning rate: 0.5

2. ROBERT options for data split (KN or RND), predict type (REG or CLAS) and hyperopt error (RMSE, etc.):

No PFI (all descriptors): PFI (only important descriptors):

split: RND split: RND type: clas type: clas error_type: acc error_type: acc



ABBREVIATIONS

Reference section for the abbreviations used.

ACC: accuracy KN: k-nearest neighbors **REG:** Regression ADAB: AdaBoost MAE: root-mean-square error RF: random forest

CSV: comma separated values MCC: Matthew's correl. coefficient RMSE: root mean square error

CLAS: classification ML: machine learning RND: random

CV: cross-validation MVL: multivariate lineal models SHAP: Shapley additive explanations

F1 score: balanced F-score NN: neural network VR: voting regressor

GB: gradient boosting PFI: permutation feature importance

GP: gaussian process R2: coefficient of determination

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This module performs RDKit conformer generation from SMILES, followed by the creation of 200+ molecular and atomic descriptors using RDKit, xTB and DBSTEP (saved as AQME-ROBERT_FILENAME.csv).

The complete output (AQME_data.dat) and raw data are stored in the AQME folder.

Time AQME: 89553.05 seconds

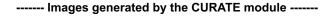


CURATE

This module takes care of data curation, including filters for correlated descriptors, noise, and duplicates, as well as conversion of categorical descriptors.

The complete output (CURATE_data.dat) and curated database are stored in the CURATE folder.

Time CURATE: 1.52 seconds





GENERATE

This module carries out a screening of ML models and selects the most accurate one. It includes a comparison of multiple hyperoptimized models and training sizes.

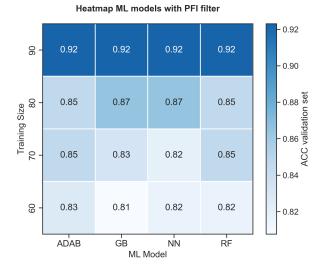
The complete output (GENERATE_data.dat) and heatmaps are stored in the GENERATE folder.

Time GENERATE: 1122.47 seconds

----- Images generated by the GENERATE module ------

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Heatmap ML models no PFI filter 0.96 8 0.96 0.88 0.94 0.92 Training Size 70 80 0.85 0.87 0.88 0.88 validation 0.88 0 0.85 0.83 0.86 0.87 0.86 0.83 0.83 0.84 0.85 9 0.84 ADAB ĠВ ŔF ΝN ML Model





Determination of predictive ability of models using four tests: 5-fold CV, y-mean (error against the mean y baseline), y-shuffle (predict with shuffled y values), and one-hot (predict using one-hot encoding instead of the X values).

The complete output (VERIFY data.dat) and donut plot are stored in the VERIFY folder.

Time VERIFY: 2.18 seconds

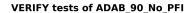
----- Images and summary generated by the VERIFY module -----

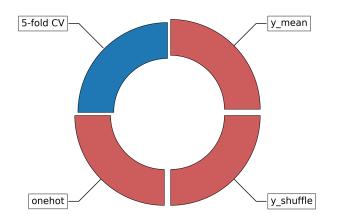
No PFI (all descriptors):

Original ACC (valid. set) 0.96 - 25% thres. = 0.72 o 5-fold CV: PASSED, ACC = 0.76, higher than thres. x y mean: FAILED, ACC = 0.92, higher than thres. x y_shuffle: FAILED, ACC = 0.81, higher than thres. x onehot: FAILED, ACC = 0.92, higher than thres.

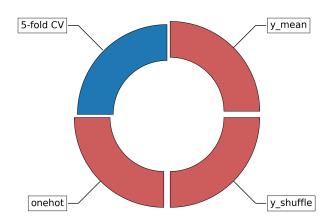
PFI (only important descriptors):

Original ACC (valid. set) 0.92 - 25% thres. = 0.69 o 5-fold CV: PASSED, ACC = 0.76, higher than thres. x y mean: FAILED, ACC = 0.92, higher than thres. x y shuffle: FAILED, ACC = 0.92, higher than thres. x onehot: FAILED, ACC = 0.92, higher than thres.





VERIFY tests of ADAB_90_PFI



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PREDICT

This module predicts and plots the results of training and validation sets from GENERATE, as well as from external test sets (if any). Feature importances from SHAP and PFI, and outlier analysis are also represented.

The complete output (PREDICT_data.dat) and heatmaps are stored in the PREDICT folder.

Time PREDICT: 15.36 seconds

----- Images and summary generated by the PREDICT module -----

No PFI (all descriptors):

Prediction metrics and descriptors

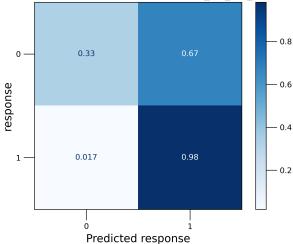
- Points Train:Validation:Test = 233:26:28
- Proportion Train:Validation:Test = 81:9:10
- Number of descriptors = 115
- Proportion (train+valid.) points:descriptors = 259:115
- Train: Accuracy = 0.83, F1 score = 0.9, MCC = 0.46
- Valid.: Accuracy = 0.96, F1 score = 0.98, MCC = 0.8
- Test: Accuracy = 0.79, F1 score = 0.88, MCC = -0.09

PFI (only important descriptors):

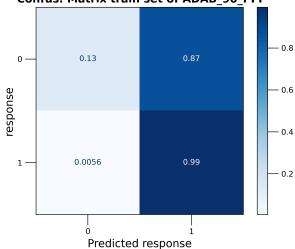
Prediction metrics and descriptors

- Points Train: Validation: Test = 233:26:28
- Proportion Train:Validation:Test = 81:9:10
- Number of descriptors = 3
- Proportion (train+valid.) points:descriptors = 259:3
- Train: Accuracy = 0.79, F1 score = 0.88, MCC = 0.28
- Valid.: Accuracy = 0.92, F1 score = 0.96, MCC = 0.0
- Test: Accuracy = 0.75, F1 score = 0.86, MCC = -0.13

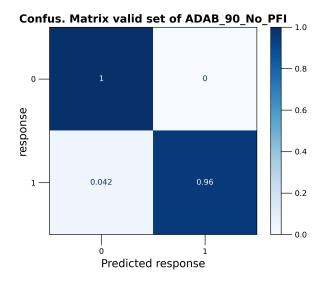


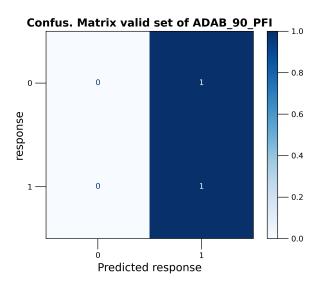


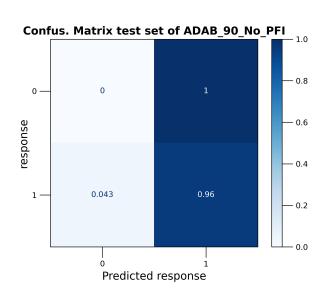
Confus. Matrix train set of ADAB 90 PFI

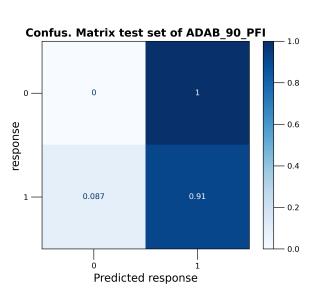


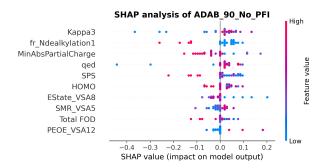
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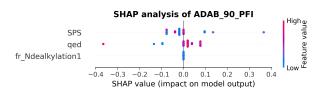






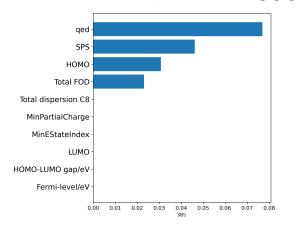




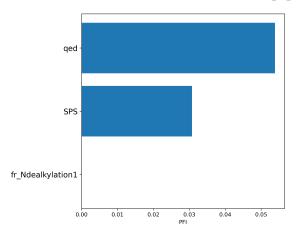


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Permutation feature importances (PFIs) of ADAB_90_No_PFI



Permutation feature importances (PFIs) of ADAB_90_PFI



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