

## Example 1: A quick look at 5 deterministic machine-learning models

Huan Tran

Some deterministic (non-probabilistic) ML models supported by matsML are introduced here. They models are

1. Support Vector Regression
2. Random Forest Regression
3. Kernel Ridge Regression
4. Gaussian Process Regression
5. Fully-Connected Neural Net

A simple dataset will be obtained from [www.matsml.org](http://www.matsml.org) for this example.

### Load data

This is a *fingerprinted* dataset, being ready for machine learning. It contains 192 compositions of hybrid organic-inorganic perovskites, each of them is represented by a fingerprint vector and the averaged band gap of multiple atomic structures predicted for this composition. This dataset was used in *Probabilistic deep learning approach for targeted hybrid organic-inorganic perovskites*, [Physical Review Materials 5, 125402 \(2021\)](#), and the raw data leading to this dataset is available at [A hybrid organic-inorganic perovskite dataset](#), [Scientific Data 4, 170057 \(2017\)](#).

```
In [1]: from matsml.data import Datasets
import pandas as pd

# obtain data
data = Datasets('fp_hoops_S1_1dest')
data.load_dataset()

# Have a look at the data fields. You will see "ID" is for the identification of the data points,
# or "fp" is the target (the averaged band gap mentioned above), and the others are the components
# of the fingerprint vector
fp_data = pd.read_csv('fp_hoops_S1_1dest.csv.gz')
print(fp_data.shape)
print(fp_data.columns)

matsML_v1.3.0
***
```

Load requested dataset(s)

Data saved in fp\_hoops\_S1\_1dest.csv.gz

(192, 34)

Index(['id', 'fp', 'mean', 'std', 'dev', 'embedding\_54', 'embedding\_116', 'embedding\_155', 'embedding\_4', 'embedding\_no', 'embedding\_136', 'embedding\_153', 'embedding\_170', 'embedding\_140', 'embedding\_179', 'embedding\_133', 'HIN4H1', 'HIN3C3', 'C4C4N4', 'H1C3C3', 'C3C3N3', 'H1C4H1', 'H1C4C4', 'H1N4O2', 'N4O2H1', 'C4C4H1', 'N4N3H1', 'H1N4N3', 'type'])

dtype='object')

\_Essential parameters of the obtained dataset, given as a dict, and needed for ML models

```
In [2]: # data parameters
data_file = 'fp_hoops_S1_1dest.csv.gz'
id_col = ['ID']
y_cols = ['mean']
comment_cols = []
n_trains = 0.9

sampling = 'random'
x_scaling = 'minmax'
y_scaling = 'normalize'

data_params = {
    'data_file': data_file,
    'id_col': id_col,
    'y_cols': y_cols,
    'comment_cols': comment_cols,
    'y_scaling': y_scaling,
    'x_scaling': x_scaling,
    'sampling': sampling,
    'n_trains': n_trains
}
```

### Model 1: Support Vector Regression

```
In [3]: from matsml.models import SVecR
# Model parameters
nfold_cv = 5
model_file = 'model_svr.pkl'
verbosity = 0
rmse_cv = False
regular_param = 2
kernel = 'rbf'
max_iter = 1

model_params = {
    'kernel': kernel,
    'nfold_cv': nfold_cv,
    'regular_param': regular_param,
    'max_iter': max_iter,
    'model_file': model_file,
    'verbosity': verbosity,
    'rmse_cv': rmse_cv
}

model = SVecR(data_params=data_params, model_params=model_params)
model.train()
model.plot(pdf_output=False)
```

Checking parameters

all passed True

Learning fingerprinted/featured data

algorithm support vector regression w/ scikit-learn

kernel rbf

regular\_param 2

max\_iter -1

nfold\_cv 5

Read data

data file fp\_hoops\_S1\_1dest.csv.gz

data size 192

training size 89.6 %

test size 10.4 %

x dimensionality 32

y dimensionality 1

y label(s) ['mean']

Scaling x minmax

xscaler saved in xscaler.pkl

Scaling y normalize

Prepare train/test sets random

Training model w/ cross validation

cv\_rmse\_train,rmse\_test,rmse\_opt: 0.0.109104 0.235800

cv\_rmse\_train,rmse\_test,rmse\_opt: 0.122489 0.180884 0.189884

cv\_rmse\_train,rmse\_test,rmse\_opt: 0.0.125387 0.172246 0.17246

cv\_rmse\_train,rmse\_test,rmse\_opt: 0.0.124511 0.164694 0.164694

cv\_rmse\_train,rmse\_test,rmse\_opt: 0.0.122357 0.261204 0.164694

SVecR model trained and saved in "model\_svr.pkl"

Now make predictions & invert scaling

unscale y: normalize Ymean 0.138838

unscale y: normalize Ymean 0.180482

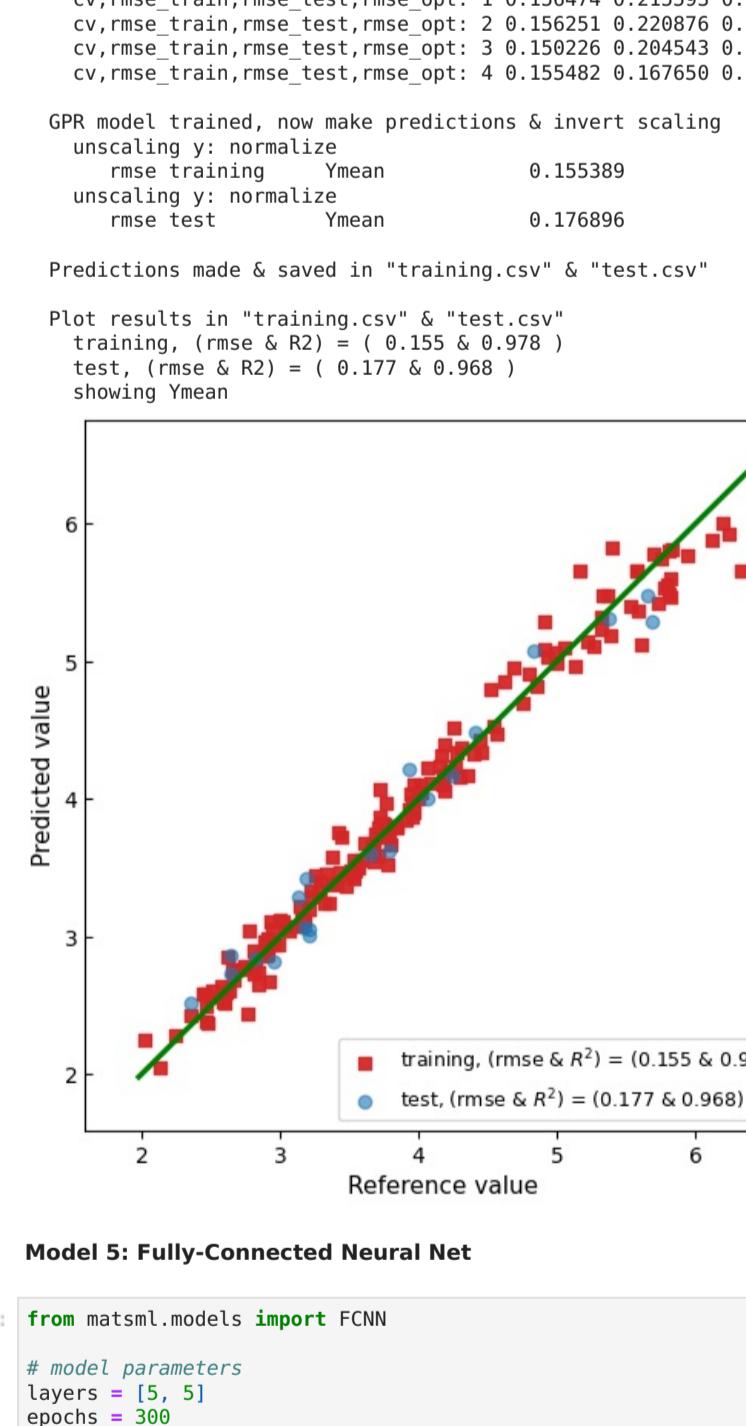
Predictions made & saved in "training.csv" & "test.csv"

Plot results in "training.csv" & "test.csv"

training, (rmse & R<sup>2</sup>) = ( 0.139 & 0.982 )

test, (rmse & R<sup>2</sup>) = ( 0.180 & 0.967 )

showing Ymean



### Model 2: Random Forest Regression

```
In [4]: from matsml.models import RFR
# Model parameters
nfold_cv = 5
model_file = 'model_rfr.pkl'
verbosity = 0
rmse_cv = False
n_estimators = 20
random_state = 11
criterion = 'friedman_mse'
max_depth = 8
get_feature_importances = True

model_params = {
    'nfold_cv': nfold_cv,
    'n_estimators': n_estimators,
    'random_state': random_state,
    'criterion': criterion,
    'max_depth': max_depth,
    'get_feature_importances': get_feature_importances,
    'model_file': model_file,
    'verbosity': verbosity,
    'rmse_cv': rmse_cv
}

model = RFR(data_params=data_params, model_params=model_params)
model.train()
model.plot(pdf_output=False)
```

Checking parameters

all passed True

Learning fingerprinted/featured data

algorithm random forest regression w/ scikit-learn

kernel rbf

nfold\_cv 5

alpha [-2, 5]

gamma [-2, 5]

n\_estimators 10

kernel = 'rbf'

model\_params = {

'kernel': kernel,

'nfold\_cv': nfold\_cv,

'model\_file': model\_file,

'alpha': alpha,

'gamma': gamma,

'n\_estimators': n\_estimators

}

Read data

data file fp\_hoops\_S1\_1dest.csv.gz

data size 192

training size 89.6 %

test size 10.4 %

x dimensionality 32

y dimensionality 1

y label(s) ['mean']

Scaling x minmax

xscaler saved in xscaler.pkl

Scaling y normalize

Prepare train/test sets random

Training model w/ cross validation

KRF model trained, now make predictions & invert scaling

unscale y: normalize Ymean 0.197194

unscale y: normalize Ymean 0.202596

Predictions made & saved in "training.csv" & "test.csv"

Plot results in "training.csv" & "test.csv"

training, (rmse & R<sup>2</sup>) = ( 0.139 & 0.982 )

test, (rmse & R<sup>2</sup>) = ( 0.207 & 0.956 )

showing Ymean



### Model 3: Kernel Ridge Regression

```
In [5]: from matsml.models import KRR
# Model parameters
nfold_cv = 5
model_file = 'model_krr.pkl'
verbosity = 0
rmse_cv = False
n_restarts_optimizer = 100
random_state = 11
criterion = 'friedman_mse'
max_depth = 8
get_feature_importances = True

model_params = {
    'nfold_cv': nfold_cv,
    'n_restarts_optimizer': n_restarts_optimizer,
    'model_file': model_file,
    'verbosity': verbosity,
    'rmse_cv': rmse_cv
}

model = KRR(data_params=data_params, model_params=model_params)
model.train()
model.plot(pdf_output=False)
```

Checking parameters

all passed True

Learning fingerprinted/featured data

algorithm kernel ridge regression w/ scikit-learn

kernel rbf

nfold\_cv 5

alpha [-2, 5]

gamma [-2, 5]

n\_restarts\_optimizer 100

random\_state 11

criterion = 'friedman\_mse'

max\_depth = 8

get\_feature\_importances True

model\_params = {

'kernel': kernel,

'nfold\_cv': nfold\_cv,

'model\_file': model\_file,

'alpha': alpha,

'gamma': gamma,

'n\_restarts\_optimizer': n\_restarts\_optimizer

'random\_state': random\_state

'criterion': criterion,

'max\_depth': max\_depth

'get\_feature\_importances': get\_feature\_importances,

'rmse\_cv': rmse\_cv

}

Read data

data file fp\_hoops\_S1\_1dest.csv.gz

data size 192

training size 89.6 %

test size 10.4 %

x dimensionality 32

y dimensionality 1

y label(s) ['mean']

Scaling x minmax

xscaler saved in xscaler.pkl

Scaling y normalize

Prepare train/test sets random

Training model w/ cross validation

KRR model trained, now make predictions & invert scaling

unscale y: normalize Ymean 0.197194

unscale y: normalize Ymean 0.202596

Predictions made & saved in "training.csv" & "test.csv"

Plot results in "training.csv" & "test.csv"

training, (rmse & R<sup>2</sup>) = ( 0.139 & 0.982 )

test, (rmse & R<sup>2</sup>) = ( 0.207 & 0.956 )

showing Ymean



### Model 4: Gaussian Process Regression

```
In [6]: from matsml.models import GPR
# Model parameters
nfold_cv = 5
model_file = 'model_gpr.pkl'
verbosity = 0
n_restarts_optimizer = 100
random_state = 11
criterion = 'mse'
max_depth = 32
get_feature_importances = True

model_params = {
    'nfold_cv
```