The raw data leading to this dataset is available at https://www.nature.com/articles/sdata201757. In [1]: from matsml.data import Datasets import pandas as pd # obtain data data=Datasets(S1='fp hoips S1 1dest') data.load dataset() # Have a look at the data fields. You will see "ID" is for the identification of the data points, # 'Ymean' is the target (the averaged band gap mentioned above), and the others are the components # of the fingeprint vector fp data = pd.read csv('fp hoips S1 ldest.csv.gz') print (fp data.shape) print (fp\_data.columns) matsML, version 1.0.0 Load requested dataset(s) Data saved in fp\_hoips\_S1\_1dest.csv.gz (192.34)Index(['Unnamed: 0', 'ID', 'Ymean', 'MagpieData avg\_dev GSvolume\_pa', 'MatscholarElementData mean embedding 54', 'MatscholarElementData std dev embedding 116', 'MatscholarElementData std dev embedding 155', 'MatscholarElementData mean embedding 4', 'PymatgenData mean mendeleev no', 'MatscholarElementData std dev embedding 136', 'MatscholarElementData std dev embedding 153', 'MatscholarElementData mean embedding 140', 'MatscholarElementData mean embedding 170', 'H1N4H1', 'H1N3H1', 'H1N3C3', 'N3C3N3', 'N3C3H1', 'H1C3C3', 'C3C3N3', 'C3N3C3', 'H1C4H1', 'H1C4C4', 'C4C4C4', 'C4C4N4', 'H1C4N4', 'C4N4H1', 'N4N3H1', 'H1N4N3', 'C4N4C4', 'H1N4O2', 'N4O2H1', 'C3C4H1', 'C4C3N3'], dtype='object') Essential parameters of the obtained dataset, given as a dict, and needed for ML models In [2]: # data parameters data file ='fp hoips S1 1dest.csv.gz' id col=['ID'] y cols=['Ymean'] comment cols=[] n trains=0.9 sampling='random' x scaling='minmax' y scaling='minmax' 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=5model 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) Learning fingerprinted/featured data support vector regression w/ scikit-learn kernel regular param max iter nfold cv Checking parameters all passed Read data data file fp hoips S1 1dest.csv.gz data size 172 (89.6 %) training size test size 20 (10.4 %) x dimensionality y dimensionality ['Ymean'] y label(s) Scaling x minmax xscaler saved in xscaler.pkl Scaling y Prepare train/test sets Training model w/ cross validation cv,rmse train,rmse test,rmse opt: 0 0.061336 0.065900 0.065900 cv,rmse\_train,rmse\_test,rmse opt: 1 0.059255 0.082091 0.065900 cv,rmse\_train,rmse\_test,rmse opt: 2 0.063549 0.065957 0.065900 cv,rmse train,rmse test,rmse opt: 3 0.061429 0.068594 0.065900 cv,rmse train,rmse test,rmse opt: 4 0.067078 0.074924 0.065900 RFR model trained and saved in "model svr.pkl" Now make predictions & invert scaling unscaling y: minmax Ymean 0.267968 rmse training unscaling y: minmax 0.303103 rmse test Ymean Predictions made & saved in "training.csv" & "test.csv" Plot results in "training.csv" & "test.csv" training, (rmse & R2) = (0.268 & 0.935)test, (rmse & R2) = (0.303 & 0.887)showing Ymean 6 5 Predicted val 3 training, (rmse &  $R^2$ ) = (0.268 & 0.935) 2 test, (rmse &  $R^2$ ) = (0.303 & 0.887) 3 Reference value **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='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) Learning fingerprinted/featured data algorithm random forest regression w/ scikit-learn nfold cv n estimators max depth criterion get feature importances random state Checking parameters all passed Read data data file fp hoips S1 1dest.csv.gz data size 192 training size 172 (89.6 %) 20 (10.4 %) test size 32 x dimensionality y dimensionality y label(s) ['Ymean'] Scaling x xscaler saved in xscaler.pkl Scaling y Prepare train/test sets Training model w/ cross validation cv,rmse train,rmse test,rmse opt: 0 0.019985 0.056001 0.056001 cv,rmse train,rmse test,rmse opt: 1 0.022022 0.049774 0.049774 cv,rmse\_train,rmse\_test,rmse\_opt: 2 0.020180 0.052332 0.049774 cv,rmse train,rmse test,rmse opt: 3 0.022245 0.055417 0.049774 cv,rmse train,rmse test,rmse opt: 4 0.020226 0.056037 0.049774 RFR model trained and saved in "model\_rfr.pkl" Top 10 features by importance MagpieData avg dev GSvolume pa importance: 0.457 MatscholarElementData mean embedding 54 importance: 0.18 MatscholarElementData std dev embedding 116 importance: 0.127 MatscholarElementData std dev embedding 155 importance: 0.074 MatscholarElementData mean embedding 4 importance: 0.044 MatscholarElementData std dev embedding 136 importance: 0.04 MatscholarElementData mean embedding 140 importance: 0.028 MatscholarElementData std dev embedding 153 importance: 0.018 MatscholarElementData mean embedding 170 importance: 0.016 PymatgenData mean mendeleev no importance: 0.004 Now make predictions & invert scaling unscaling y: minmax rmse training Ymean 0.128367 unscaling y: minmax rmse test Ymean 0.2225 Predictions made & saved in "training.csv" & "test.csv" Plot results in "training.csv" & "test.csv" training, (rmse & R2) = (0.128 & 0.984)test, (rmse & R2) = (0.222 & 0.961)showing Ymean 6 5 Predicted value 3 training, (rmse &  $R^2$ ) = (0.128 & 0.984) 2 test, (rmse &  $R^2$ ) = (0.222 & 0.961) 3 Reference value Model 3: Kernel Ridge Regression (KRR) In [5]: from matsml.models import KRR # Model parameters nfold cv = 5model file = 'model krr.pkl' alpha = [-2, 5]gamma = [-2, 5]n grids = 10kernel = 'rbf' model params={'kernel':kernel,'nfold cv':nfold cv,'model file':model file,'alpha':alpha, 'gamma':gamma,'n grids':n grids} model = KRR(data params=data params, model params=model params) model.train() model.plot(pdf output=False) Learning fingerprinted/featured data algorithm kernel ridge regression w/ scikit-learn kernel nfold cv [-2, 5]alpha gamma [-2, 5]number of alpha/gamma grids Checking parameters all passed Read data fp hoips S1 1dest.csv.gz data file data size 192 training size 172 (89.6 %) 20 (10.4 %) test size x dimensionality 32 y dimensionality y label(s) ['Ymean'] Scaling x minmax xscaler saved in xscaler.pkl Scaling y minmax Prepare train/test sets random Building model Training model w/ cross validation KRR model trained, now make predictions & invert scaling unscaling y: minmax 0.200223 rmse training Ymean unscaling y: minmax 0.154706 rmse test Ymean Predictions made & saved in "training.csv" & "test.csv" Plot results in "training.csv" & "test.csv" training, (rmse & R2) = (0.200 & 0.964)test, (rmse & R2) = (0.155 & 0.973)showing Ymean 6 5 Predicted value 3 training, (rmse &  $R^2$ ) = (0.200 & 0.964) 2 test, (rmse &  $R^2$ ) = (0.155 & 0.973) 2 3 6 5 Reference value **Model 4: Gaussian Process Regression** In [6]: from matsml.models import GPR # Model parameters nfold cv=5model\_file='model\_gpr.pkl' verbosity=0 rmse cv=True n\_restarts\_optimizer=100 model params={'nfold cv':nfold cv,'n restarts optimizer':n restarts optimizer,'model file':model file, 'verbosity':verbosity,'rmse cv':rmse cv} model=GPR(data params=data params, model params=model params) model.train() model.plot(pdf output=False) Learning fingerprinted/featured data gaussian process regression w/ scikit-learn algorithm nfold cv fmin l bfgs b optimizer n\_restarts\_optimizer True rmse cv Checking parameters True all passed Read data data file fp\_hoips\_S1\_1dest.csv.gz data size 192 172 (89.6 %) training size 20 (10.4 %) test size 32 x dimensionality y dimensionality 1 ['Ymean'] y label(s) Scaling x minmax xscaler.pkl xscaler saved in Scaling y minmax Prepare train/test sets Training model w/ cross validation cv,rmse\_train,rmse\_test,rmse\_opt: 0 0.017592 0.036452 0.036452 unscaling y: minmax rmse cv\_test Ymean 0.156809 cv,rmse\_train,rmse\_test,rmse\_opt: 1 0.018102 0.030700 0.030700 unscaling y: minmax rmse cv test Ymean 0.132068 cv,rmse\_train,rmse\_test,rmse\_opt: 2 0.015147 0.032244 0.030700 unscaling y: minmax rmse cv\_test Ymean 0.138708 cv,rmse\_train,rmse\_test,rmse\_opt: 3 0.018131 0.049642 0.030700 unscaling y: minmax rmse cv test 0.213553 Ymean cv,rmse\_train,rmse\_test,rmse\_opt: 4 0.018877 0.031716 0.030700 unscaling y: minmax rmse cv\_test Ymean 0.136439 GPR model trained, now make predictions & invert scaling unscaling y: minmax 0.079571 rmse training Ymean unscaling y: minmax rmse test Ymean 0.154774 Predictions made & saved in "training.csv" & "test.csv" Plot results in "training.csv" & "test.csv" training, (rmse & R2) = (0.080 & 0.994)test, (rmse & R2) = (0.155 & 0.983)showing Ymean 6

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

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.

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

models

**Huan Tran** 

Load data

Support Vector Regression
Random Forest Regression
Kernel Ridge Regression

4. Gaussian Process Regression5. Fully-Connected Neural Net

A simple dataset will be obtained from www.matsml.org for this example.

Predi 3 training, (rmse &  $R^2$ ) = (0.080 & 0.994) 2 test, (rmse &  $R^2$ ) = (0.155 & 0.983) 3 Reference value **Model 5: Fully-Connected Neural Net** In [7]: from matsml.models import FCNN # model parameters layers=[5,5] epochs=300 nfold cv=5 use bias=True model\_file='model\_fcnn.pkl' loss='mse' verbosity=0 batch size=32 activ funct='elu' optimizer='nadam' model\_params={'layers':layers,'activ\_funct':activ\_funct,'epochs':epochs,'nfold\_cv':nfold\_cv, 'optimizer':optimizer, 'use\_bias':use\_bias, 'model\_file':model\_file, 'loss':loss, 'batch\_size':batch\_size,'verbosity':verbosity,'rmse\_cv':False} model=FCNN(data\_params=data\_params, model\_params=model\_params) model.train() model.plot(pdf\_output=False) Learning fingerprinted/featured data fully connected NeuralNet w/ TensorFlow algorithm layers [5, 5] activ funct elu 300 epochs optimizer nadam nfold cv Checking parameters all passed True Read data data file fp\_hoips\_S1\_1dest.csv.gz data size 192 172 (89.6 %) training size test size 20 (10.4 %) x dimensionality y dimensionality y label(s) ['Ymean'] Scaling x minmax xscaler saved in xscaler.pkl Scaling y minmax Prepare train/test sets Building model Training model w/ cross validation cv,rmse train,rmse test,rmse opt: 0 0.053168 0.066369 0.066369 cv,rmse\_train,rmse\_test,rmse\_opt: 1 0.037757 0.042781 0.042781 cv,rmse\_train,rmse\_test,rmse\_opt: 2 0.031396 0.043164 0.042781 cv,rmse\_train,rmse\_test,rmse\_opt: 3 0.029349 0.040847 0.040847 cv,rmse\_train,rmse\_test,rmse\_opt: 4 0.028165 0.042264 0.040847 Optimal ncv: 3 ; optimal NET saved FCNN trained, now make predictions & invert scaling unscaling y: minmax 0.137451 Ymean rmse training unscaling y: minmax rmse test Ymean 0.130711 Predictions made & saved in "training.csv" & "test.csv" Plot results in "training.csv" & "test.csv" training, (rmse & R2) = (0.137 & 0.983)test, (rmse & R2) = (0.131 & 0.979)showing Ymean 6

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training, (rmse &  $R^2$ ) = (0.137 & 0.983)

6

test, (rmse &  $R^2$ ) = (0.131 & 0.979)

Reference value

Predicted value