	<ul> <li>5. Fully-Connected Neural Net</li> <li>A simple dataset will be obtained from www.matsml.org for this example.</li> <li>Load data</li> <li>This is a <i>fingerprinted</i> dataset, being ready for machine learning. It contains 192 compositions of hybrid organic-inorganic perovskites,</li> </ul>
[1]:	each of them is represented by a fingerprint vector and the averaged band gap of multiple atomic structures predicted for this composition. The raw data leading to this dataset is available at <a href="https://www.nature.com/articles/sdata201757">https://www.nature.com/articles/sdata201757</a> . <b>from</b> matsml.data <b>import</b> Datasets <b>import</b> pandas <b>as</b> 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_1dest.csv.gz')  print (fp_data.shape)  print (fp_data.columns)  matsML, v1.0.1  *****
	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',
[2]:	'MatscholarElementData mean 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
[2].	<pre># 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,</pre>
[3]:	<pre>Model 1: Support Vector Regression  from matsml.models import SVecR  # Model parameters nfold_cv=5 model_file='model_svr.pkl' verbosity=0</pre>
	<pre>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)</pre>
	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_hoips_S1_1dest.csv.gz data size 192 training size 172 (89.6 %) test size 20 (10.4 %) x dimensionality 32 y dimensionality 1 y label(s) ['Ymean'] Scaling x minmax xscaler saved in xscaler.pkl
	Scaling y minmax Prepare train/test sets random Training model w/ cross validation    cv,rmse_train,rmse_test,rmse_opt: 0 0.062320 0.078931 0.078931    cv,rmse_train,rmse_test,rmse_opt: 1 0.067054 0.072070 0.072070    cv,rmse_train,rmse_test,rmse_opt: 2 0.066908 0.073685 0.072070    cv,rmse_train,rmse_test,rmse_opt: 3 0.062854 0.077470 0.072070    cv,rmse_train,rmse_test,rmse_opt: 4 0.062729 0.061967 0.061967 RFR model trained and saved in "model_svr.pkl" Now make predictions & invert scaling    unscaling y: minmax
	rmse training Ymean 0.269206 unscaling y: minmax rmse test Ymean 0.285963 Predictions made & saved in "training.csv" & "test.csv" Plot results in "training.csv" & "test.csv" training, (rmse & R2) = (0.269 & 0.933) test, (rmse & R2) = (0.286 & 0.929) showing Ymean
	Predicted value
	training, (rmse & $R^2$ ) = (0.269 & 0.933) test, (rmse & $R^2$ ) = (0.286 & 0.929) Reference value
[4]:	<pre>Model 2: Random Forest Regression  from matsml.models import RFR  # Model parameters nfold_cv=5 model_file='model_rfr.pkl' verbosity=0</pre>
	<pre>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}</pre>
	<pre>model=RFR(data_params=data_params, model_params=model_params) model.train() model.plot(pdf_output=False)  Checking parameters    all passed</pre>
	nfold_cv 5 n_estimators 20 max_depth 8 criterion mse get_feature_importances True random_state 11 Read data data file fp_hoips_S1_1dest.csv.gz data size 192 training size 172 (89.6 %)
	test size 20 (10.4 %)  x dimensionality 32  y dimensionality 1  y label(s) ['Ymean']  Scaling x minmax  xscaler saved in xscaler.pkl  Scaling y minmax  Prepare train/test sets random  Training model w/ cross validation  cv, rmse_train, rmse_test, rmse_opt: 0 0.022399 0.046180 0.046180
	cv,rmse_train,rmse_test,rmse_opt: 1 0.022101 0.046171 0.046171 cv,rmse_train,rmse_test,rmse_opt: 2 0.022216 0.062874 0.046171 cv,rmse_train,rmse_test,rmse_opt: 3 0.019279 0.061148 0.046171 cv,rmse_train,rmse_test,rmse_opt: 4 0.023358 0.054302 0.046171 RFR model trained and saved in "model_rfr.pkl" Top 10 features by importance MatscholarElementData std_dev embedding 116 importance: 0.426 MagpieData avg_dev GSvolume_pa importance: 0.214 MatscholarElementData std_dev embedding 136 importance: 0.095 MatscholarElementData mean embedding 4 importance: 0.079 MatscholarElementData mean embedding 54 importance: 0.063
	MatscholarElementData std_dev embedding 155 importance: 0.042 MatscholarElementData mean embedding 170 importance: 0.025 MatscholarElementData std_dev embedding 153 importance: 0.022 MatscholarElementData mean embedding 140 importance: 0.017 PymatgenData mean mendeleev_no importance: 0.004 Now make predictions & invert scaling unscaling y: minmax rmse training Ymean 0.123401 unscaling y: minmax rmse test Ymean 0.186584
	Predictions made & saved in "training.csv" & "test.csv"  Plot results in "training.csv" & "test.csv"  training, (rmse & R2) = (0.123 & 0.986)  test, (rmse & R2) = (0.187 & 0.952)  showing Ymean
	Predicted value  4  3
	training, (rmse & $R^2$ ) = (0.123 & 0.986) test, (rmse & $R^2$ ) = (0.187 & 0.952)  Reference value  Model 3: Kernel Ridge Regression (KRR)
[5]:	<pre>from matsml.models import KRR  # Model parameters nfold_cv = 5 model_file = 'model_krr.pkl' alpha = [-2,5] gamma = [-2,5] n_grids = 10 kernel = 'rbf'</pre>
	<pre>model_params={'kernel':kernel,'nfold_cv':nfold_cv,'model_file':model_file,'alpha':alpha,</pre>
	Learning fingerprinted/featured data  algorithm kernel ridge regression w/ scikit-learn  kernel rbf  nfold_cv 5  alpha [-2, 5]  gamma [-2, 5]  number of alpha/gamma grids 10  Read data  data file fp_hoips_S1_1dest.csv.gz  data size 192
	training size 172 (89.6 %)  test size 20 (10.4 %)  x dimensionality 32  y dimensionality 1  y label(s) ['Ymean']  Scaling x minmax  xscaler saved in xscaler.pkl  Scaling y minmax  Prepare train/test sets random  Building model KRR
	Training model w/ cross validation  KRR model trained, now make predictions & invert scaling    unscaling y: minmax    rmse training
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	$\frac{1}{3}$ training, (rmse & $R^2$ ) = (0.186 & 0.967) test, (rmse & $R^2$ ) = (0.317 & 0.920)
[6]:	Model 4: Gaussian Process Regression  from matsml.models import GPR  # Model parameters  nfold even
	<pre>nfold_cv=5 model_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,</pre>
	model.plot(pdf_output=False)  Checking parameters all passed True  Learning fingerprinted/featured data algorithm gaussian process regression w/ scikit-learn nfold_cv 5 optimizer fmin_l_bfgs_b
	n_restarts_optimizer 100 rmse_cv True  Read data data file fp_hoips_S1_1dest.csv.gz data size 192 training size 172 (89.6 %) test size 20 (10.4 %) x dimensionality 32 y dimensionality 1 y label(s) ['Ymean']
	Scaling x minmax xscaler saved in xscaler.pkl Scaling y minmax Prepare train/test sets random Training model w/ cross validation cv,rmse_train,rmse_test,rmse_opt: 0 0.020118 0.034120 0.034120 unscaling y: minmax rmse cv_test Ymean 0.146779 cv,rmse_train,rmse_test,rmse_opt: 1 0.016859 0.040843 0.034120 unscaling y: minmax rmse cv_test Ymean 0.175702
	<pre>cv,rmse_train,rmse_test,rmse_opt: 2 0.019410 0.038437 0.034120 unscaling y: minmax     rmse cv_test</pre>
	<pre>unscaling y: minmax     rmse training</pre>
	Predicted value
	training, (rmse & $R^2$ ) = (0.087 & 0.993) test, (rmse & $R^2$ ) = (0.097 & 0.991) Reference value
[7]:	Model 5: Fully-Connected Neural Net  from matsml.models import FCNN  # model parameters layers=[5,5] epochs=300 nfold_cv=5
	<pre>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,</pre>
	<pre>'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)  Checking parameters all passed True</pre>
	Learning fingerprinted/featured data algorithm fully connected NeuralNet w/ TensorFlow layers [5, 5] activ_funct elu epochs 300 optimizer nadam nfold_cv 5 Read data data file fp_hoips_S1_1dest.csv.gz data size 192 training size 172 (89.6 %)
	test size 20 (10.4 %)  x dimensionality 32  y dimensionality 1  y label(s) ['Ymean']  Scaling x minmax  xscaler saved in xscaler.pkl  Scaling y minmax  Prepare train/test sets random  Building model FCNN  Training model w/ cross validation
	<pre>cv,rmse_train,rmse_test,rmse_opt: 0 0.046565 0.058864 0.058864 cv,rmse_train,rmse_test,rmse_opt: 1 0.032680 0.040572 0.040572 cv,rmse_train,rmse_test,rmse_opt: 2 0.028597 0.041473 0.040572 cv,rmse_train,rmse_test,rmse_opt: 3 0.027411 0.041591 0.040572 cv,rmse_train,rmse_test,rmse_opt: 4 0.026858 0.033028 0.033028 Optimal ncv: 4 ; optimal NET saved FCNN trained, now make predictions &amp; invert scaling</pre>
	unscaling y: minmax rmse training Ymean 0.121247 unscaling y: minmax
	rmse training Ymean 0.121247