## First 20 Models Simultaneous Run

**General Imports** 

```
import pandas as pd
import os
import matplotlib.pyplot as plt
import numpy as np
from sklearn.model_selection import StratifiedShuffleSplit
from sklearn.preprocessing import StandardScaler
from sklearn.model_selection import cross_validate
from sklearn.model_selection import KFold

from sklearn.utils import all_estimators
```

**Automatically Importing All Regressions** 

```
In []: estimators = all_estimators(type_filter='regressor')

all_regs = []
for name, RegressorClass in estimators:
    try:
        if name != 'DummyRegressor' and name != 'GaussianProcessRegressor':
            print('Appending', name)
            reg = RegressorClass()
            all_regs.append(reg)
    except Exception as e:
        print(e)
```

```
Appending ARDRegression
Appending AdaBoostRegressor
Appending BaggingRegressor
Appending BayesianRidge
Appending CCA
Appending DecisionTreeRegressor
Appending ElasticNet
Appending ElasticNetCV
Appending ExtraTreeRegressor
Appending ExtraTreesRegressor
Appending GammaRegressor
Appending GradientBoostingRegressor
Appending HistGradientBoostingRegressor
Appending HuberRegressor
Appending IsotonicRegression
Appending KNeighborsRegressor
Appending KernelRidge
Appending Lars
Appending LarsCV
Appending Lasso
Appending LassoCV
Appending LassoLars
Appending LassoLarsCV
Appending LassoLarsIC
Appending LinearRegression
Appending LinearSVR
Appending MLPRegressor
Appending MultiOutputRegressor
init () missing 1 required positional argument: 'estimator'
Appending MultiTaskElasticNet
Appending MultiTaskElasticNetCV
Appending MultiTaskLasso
Appending MultiTaskLassoCV
Appending NuSVR
Appending OrthogonalMatchingPursuit
Appending OrthogonalMatchingPursuitCV
Appending PLSCanonical
Appending PLSRegression
Appending PassiveAggressiveRegressor
Appending PoissonRegressor
Appending QuantileRegressor
Appending RANSACRegressor
Appending RadiusNeighborsRegressor
Appending RandomForestRegressor
Appending RegressorChain
init () missing 1 required positional argument: 'base estimator'
Appending Ridge
Appending RidgeCV
Appending SGDRegressor
Appending SVR
Appending StackingRegressor
init () missing 1 required positional argument: 'estimators'
Appending TheilSenRegressor
Appending TransformedTargetRegressor
Appending TweedieRegressor
Appending VotingRegressor
init () missing 1 required positional argument: 'estimators'
```

```
In [ ]: def load_pp_data():
            csv_path = r"C:\Users\18123\OneDrive\Documents\IUBloomington\Machine-Learning-Proj
            return pd.read csv(csv path)
        pp = load_pp_data()
        print(pp.describe())
                                                                           PΕ
                        ΑT
                                     V
                                                 AΡ
                                                              RH
        count 9568.000000 9568.000000 9568.000000 9568.000000 9568.000000
        mean
                 19.651231
                             54.305804 1013.259078
                                                       73.308978
                                                                  454.365009
        std
                  7.452473 12.707893
                                           5.938784
                                                       14.600269
                                                                   17.066995
                 1.810000
                              25.360000 992.890000
        min
                                                       25.560000
                                                                  420.260000
        25%
                 13.510000
                             41.740000 1009.100000
                                                       63.327500
                                                                   439.750000
        50%
                 20.345000
                              52.080000 1012.940000
                                                       74.975000
                                                                  451.550000
        75%
                 25.720000
                              66.540000 1017.260000
                                                       84.830000
                                                                   468.430000
                 37.110000
                             81.560000 1033.300000
                                                      100.160000
                                                                  495.760000
        max
        Train/Test Split and Preprocess Data
In []: pp["AT cat"] = pd.cut(pp["AT"],bins=[0.,10.,20.,30.,np.inf],labels=[1,2,3,4])
        split = StratifiedShuffleSplit(n splits=1,test size=0.2,random state=42)
        for train_index, test_index in split.split(pp,pp["AT_cat"]):
            train_set = pp.loc[train_index]
            test set = pp.loc[test index]
        for set_ in(train_set,test_set):
            set_.drop("AT_cat",axis=1,inplace=True)
        pptrain = train set.copy()
        pptest = test_set.copy()
        pptrain_attrib = pptrain.drop("PE",axis=1)
        pptrain_labels = pptrain["PE"].copy()
        pptest_attrib = pptest.drop("PE",axis=1)
        pptest_labels = pptest["PE"].copy()
        scaler = StandardScaler()
        scaler.fit_transform(pptrain_attrib)
Out[]: array([[ 1.1978498 , 0.96554795, 0.37377565, -2.67409022],
               [0.64009018, -1.03750958, -1.88469509, -2.35340963],
               [-1.82211612, -1.45609422, -0.36887464, 1.17611946],
               . . . ,
               [-1.07754063, -0.84989538, 0.57724148, 0.20454577],
               [-0.67971691, -0.96104497, 0.78748951, 0.87314098],
               [ 0.89545 , 0.56351752, -0.13658448, -1.12107019]])
        Simultaneous Run
In [ ]:
        def run(model):
            print(f"checking {model}")
            try:
                cv_outer = KFold(n_splits=10, shuffle=True, random_state=2)
```

```
cv_output_dict = cross_validate(model, pptrain_attrib, pptrain_labels, scoring
        return cv_output_dict
    except:
        pass
def comparison(modellst):
    cv_data = []
    errors = []
    passed_models = []
    for i in range(len(modellst)):
        x = run(modellst[i])
        if type(x) == dict:
            cv_data += [x]
        else:
            errors += [i]
    for j in range(len(modellst)):
        if j not in errors:
            passed_models += [modellst[j]]
    return vizualize(cv_data, passed_models)
def vizualize(cv_data, modellst):
    return box_rmse(cv_data, modellst, 'train'), box_rmse(cv_data, modellst, 'test'),
def runtime(cv_data, modellst):
    timefig = plt.figure(constrained_layout=True)
    df = pd.DataFrame()
    for i,j in zip(cv_data,modellst):
        df[j] = list(i[('fit_time')])
    sorted index = df.median().sort values(ascending=False).index
    df_sorted=df[sorted_index]
    # sns.boxplot(data=df_sorted, orient='h')
    df_sorted.boxplot(vert=False,grid=False,color='purple')
    plt.xlabel('Run Time')
    plt.ylabel('Models')
    return timefig
def box_rmse(cv_data, modellst, data_split):
    RMSEfig = plt.figure(constrained_layout=True)
    df = pd.DataFrame()
    for i, j in zip(cv data, modellst):
        df[j] = list(np.sqrt(i[data_split+'_neg_mean_squared_error']*-1))
    sorted_index = df.median().sort_values(ascending=False).index
    df sorted=df[sorted index]
    # sns.boxplot(data=df_sorted, orient='h')
    df_sorted.boxplot(vert=False,grid=False,color='purple')
    plt.xlabel(f'{data_split} Root Mean Squared Error (Lower is better)')
    return RMSEfig
def box_r2(cv_data, modellst, data_split):
    R2fig = plt.figure(constrained_layout=True)
    df = pd.DataFrame()
    for i,j in zip(cv_data,modellst):
        df[j] = list(i[data_split+'_r2'])
    sorted_index = df.median().sort_values().index
    df_sorted=df[sorted_index]
    # sns.boxplot(data=df sorted, orient='h')
```

```
df sorted.boxplot(vert=False,grid=False,color='purple')
    plt.xlabel(f'{data_split} R-Squared Score (Higher is better)')
    return R2fig
def box mae(cv data, modellst, data split):
    MAEfig = plt.figure(constrained_layout=True)
    df = pd.DataFrame()
    for i,j in zip(cv_data,modellst):
        df[j] = list(i[data_split+'_neg_mean_absolute_error']*-1)
    sorted_index = df.median().sort_values(ascending=False).index
    df sorted=df[sorted index]
    # sns.boxplot(data=df_sorted, orient='h')
    df sorted.boxplot(vert=False,grid=False,color='purple')
    plt.xlabel(f'{data_split} Mean Absolute Error (Lower is better)')
    return MAEfig
y = all_regs[0:20]
x = all_regs[0:3]
comparison(y)
plt.show()
checking ARDRegression()
checking AdaBoostRegressor()
checking BaggingRegressor()
checking BayesianRidge()
checking CCA()
checking DecisionTreeRegressor()
checking ElasticNet()
checking ElasticNetCV()
checking ExtraTreeRegressor()
checking ExtraTreesRegressor()
checking GammaRegressor()
checking GradientBoostingRegressor()
checking HistGradientBoostingRegressor()
checking HuberRegressor()
c:\Users\18123\AppData\Local\Programs\Python\Python38\lib\site-packages\sklearn\linea
r_model\_huber.py:335: ConvergenceWarning: lbfgs failed to converge (status=1):
STOP: TOTAL NO. of ITERATIONS REACHED LIMIT.
Increase the number of iterations (max_iter) or scale the data as shown in:
    https://scikit-learn.org/stable/modules/preprocessing.html
  self.n_iter_ = _check_optimize_result("lbfgs", opt_res, self.max_iter)
checking IsotonicRegression()
checking KNeighborsRegressor()
checking KernelRidge()
checking Lars()
checking LarsCV()
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```
c:\Users\18123\AppData\Local\Programs\Python\Python38\lib\site-packages\sklearn\linea
r model\ base.py:133: FutureWarning: The default of 'normalize' will be set to False
in version 1.2 and deprecated in version 1.4.
If you wish to scale the data, use Pipeline with a StandardScaler in a preprocessing
stage. To reproduce the previous behavior:
from sklearn.pipeline import make pipeline
model = make_pipeline(StandardScaler(with_mean=False), Lars())
If you wish to pass a sample_weight parameter, you need to pass it as a fit parameter
to each step of the pipeline as follows:
kwargs = \{s[0] + ' \text{ sample weight': sample weight for s in model.steps}\}
model.fit(X, y, **kwargs)
 warnings.warn(
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model = make_pipeline(StandardScaler(with_mean=False), LarsCV())

If you wish to pass a sample_weight parameter, you need to pass it as a fit parameter
to each step of the pipeline as follows:

kwargs = {s[0] + '__sample_weight': sample_weight for s in model.steps}
model.fit(X, y, **kwargs)

warnings.warn(
checking Lasso()
```







