a:	Decision Tree Introduction We will be using the wine quality data set for these exercises. This data set contains various chemical properties of wine, such as acidity, sugar, pH, and alcohol. It also contains a quality metric (3-9, with highest being better) and a color (red or white). The name of the file is Wine_Quality_Data.csv.						
	 import os, pandas as Import the data and ex We will be using all of t ### BEGIN SOLUTION 	pd, numpy a amine the featu	e file is Wine_Qu s np, matplot1 ures. color (white o				
]:	<pre>filepath = 'Wine_Qua' data = pd.read_csv("d data.head() fixed_acidity volatile_ac 0 7.4</pre>	C:\\Users\\r	d residual_sugar				
: 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	2 7.83 11.2	0.76 0.0 0.28 0.5 0.70 0.0 float64 float64 float64 float64 float64 float64 float64 float64 float64 float64 float64 float64 object	2.3 66 1.9	0.092 15.0 54.0 0.9970 3.26 0.65 9.8 5 red 0.075 17.0 60.0 0.9980 3.16 0.58 9.8 6 red			
C:	dtype: object Convert the color feature to data['color'] = data ### END SOLUTION • Use StratifiedShu	an integer. The color repla	ce('white',0).	replace('red',1).astype(np.int) ain and test sets that are stratified by wine quality. If possible, preserve the indices of the split below.			
: [<pre>### BEGIN SOLUTION # All data columns ex feature_cols = [x for from sklearn.model_se</pre>	xcept for co r x in data. election imp	<i>lor</i> columns if x n ort Stratified	ShuffleSplit			
N	<pre># Get the index value train_idx, test_idx: # Create the data se X_train = data.loc[tr y_train = data.loc[tr X_test = data.loc[test y_test = data.loc[test y_test = data.loc[test y_test = data.loc[test Now check the percent come.</pre>	erator StratifiedSh es from the next(strat ts rain_idx, fe rain_idx, 'c st_idx, feat st_idx, 'col nposition of each	uffleSplit(n_s generator _shuff_split.s ature_cols] olor'] ure_cols] or'] ch quality level in t	splits=1, test_size=1000, random_state=42) split(data[feature_cols], data['color'])) the train and test data sets. The data set is mostly white wine, as can be seen below.			
: 0 1 N	y_train.value_counts	(normalize= T loat64	rue).sort_inde	ex()			
ľ	1 0.246Name: color, dtype: fFitting a decision tree cDetermine how many r	classifier with n	ent and what the c	ximum depth, features, or leaves. depth of this (very large) tree is. In and test data sets. What do you think is going on here based on the differences in prediction error?			
T :	<pre>from sklearn.tree im dt = DecisionTreeClas dt = dt.fit(X_train, The number of nodes and t # without pruning dt.treenode_count,</pre>	ssifier(rand y_train) he maximum a	om_state=42) ctual depth.				
A :	<pre>def measure_error(y_</pre>	<pre>import accu true, y_pred ({'accuracy'</pre>	, label): :accuracy_scor	recision_score, recall_score, f1_score re(y_true, y_pred), score(y_true, y_pred),			
:	<pre>def measure_error(y_ return pd.Series</pre>	<pre>'recall': 'f1': f1_s name=label true, y_pred ({'accuracy' 'precision 'recall':</pre>	<pre>recall_score(y core(y_true, y) ,train): :accuracy_scor ': precision_s</pre>	<pre>/_true, y_pred), /_pred)}, re(y_true, y_pred), recore(y_true, y_pred), /_true, y_pred),</pre>			
:	<pre>def measure_error(y_ return pd.Series</pre>	re(y_true, y_pred), score(y_true, y_pred), y_true, y_pred), y_pred)},					
e	# The error on the ty_train_pred = dt.predy_test_pred = dt.pred	n here. raining and edict(X_traidict(X_test)) r = pd.conca	test data sets n) t([measure_err	than the test data, which is consistent with (mild) overfitting. Also notice the perfect recall score for the training data. In many instances, this prediction difference is for(y_train, y_train_pred, 'train'), test, y_test_pred, 'test')],			
: _	train_test_full_error ### END SOLUTION train t accuracy 0.999818 0.9840	ax r est	is=1) # we con	catenate these two together, re saying we're concantenating across the columns.			
	recall 1.000000 0.995145 f1 0.999631 0.9967611 • Using grid search with cross validation, find a decision tree that performs well on the test data set. Use a different variable name for this decision tree model • Determine the number of nodes and the depth of this tree. • Measure the errors on the training and test sets as before and compare them to those from the tree.						
w th	• Measure the errors on the training and test sets as before and compare them to those from the tree . we use max_depth as one of the hyperparameters and max_features as one of the parameters as well. So we have to make sure that those names match up. We pass in a list of values. So here it's going to be a range of one through the max_depth that we calculated, plus one so that we're able to include the last one counting by two, and then this is also going to be using the decision tree we classified, and the number of features that were used there. Generally, this is going to be all of the features if we didn't prune at all. ### BEGIN SOLUTION from sklearn.model_selection import GridSearchCV param_grid = {'max_depth':range(1, dt.treemax_depth+1, 2),						
: (GR.best_estimatort. # smaller tree compa	reenode_co red to earli	unt, GR.best_e er(171,22)	estimatortreemax_depth So it would seem the previous example overfit the data, but only slightly so.			
:	<pre>y_train_pred_gr = GR y_test_pred_gr = GR. train_test_gr_error train_test_gr_error #most of them we see</pre>	<pre>.predict(X_t predict(X_te = pd.concat(improvement</pre>	rain) st) [measure_error measure_error axis=1) s in scores	(y_train, y_train_pred_gr, 'train'), (y_test, y_test_pred_gr, 'test')],			
:	<pre># not much improvement for accuracy # improvement in precision. # same for recall #improvement in our f1 score. train test accuracy 0.995816 0.989000 precision 0.998501 0.983539</pre>						
	recall 0.984479 0.9718 f1 0.991440 0.9778 Re-split the data into > the data. Using grid search with Make a plot of actual v. ### BEGIN SOLUTION feature_cols = [x for the data se	the data into X and y parts, this time with residual_sugar being the predicted (y) data. <i>Note:</i> if the indices were preserved from the StratifiedShuffleSplit output, they can be used again to split a. rid search with cross validation, find a decision tree regression model that performs well on the test data set. *since this is regression we measure the errors on the training and test sets using mean squared error plot of actual vs predicted residual sugar. N SOLUTION cols = [x for x in data.columns if x != 'residual_sugar'] the data sets					
	#We are then going to set x_train equal to our original data using #train index that we defined earlier using stratified shuffle split. X_train = data.loc[train_idx, feature_cols] y_train = data.loc[train_idx, 'residual_sugar'] X_test = data.loc[test_idx, feature_cols] y_test = data.loc[test_idx, 'residual_sugar'] from sklearn.tree import DecisionTreeRegressor						
	r = DecisionTreeRegressor().fit(X_train, y_train) aram_grid = {'max_depth':range(1, dr.treemax_depth+1, 2),						
T :	n_jobs=-1) #So in reality we are minimizing mean squared error GR_sugar = GR_sugar.fit(X_train, y_train) The number of nodes and the maximum depth of the tree. This tree has lots of nodes, which is not so surprising given the continuous data. GR_sugar.best_estimatortreenode_count, GR_sugar.best_estimatortreemax_depth # lot more increased in nodes , and depth is 13 compared to earlier (2891, 13)						
:	from sklearn.metrics #we can't use the sat #we use our GR_ sugat #on the set that we y_train_pred_gr_sugar y_test_pred_gr_sugar	<pre>import mean me classific r to predict trained on, r = GR_sugar = GR_sugar</pre>	_squared_error ation matrix t our actual x_ as well as the .predict(X_tra .predict(X_tes	that we just used before. train while we would predict prediction on our x_test. in) it)			
: _	<pre>train_test_gr_sugar_error = pd.Series({'train': mean_squared_error(y_train, y_train_pred_gr_sugar),</pre>						
:	A plot of actual vs predicted residual sugar. sns.set_context('notebook') sns.set_style('white') fig = plt.figure(figsize=(6,6)) ax = plt.axes() ph_test_predict = pd.DataFrame({'test':y_test.values,						
	25 predict 10 5						
-	-c conda-forge pydot	plus) can be	orogram (GraphViz	z) and Python library (PyDotPlus). GraphViz can be installed with a package manager on Linux and Mac. For PyDotPlus, either pip or conda (conda instal library.			
T:	 Creating a visualization The decision tree from this 	n of the decision of the decision will have too mel to use n	n tree where wine any nodes to visu	e color was predicted and the number of features and/or splits are not limited. e color was predicted but a grid search was used to find the optimal depth and number of features. palize. and packages, variables will be lost			
() ; ; ; ; ; ; ; ; ; ; ; ; ; ; ; ; ; ;	Collecting Graphviz Downloading graphvi Installing collected Successfully installe Note: you may need to #install pydotplus conda install -c cond	z-0.19.1-py3 packages: Gr d Graphviz-6 restart the da-forge pyd	aphviz 1.19.1 kernel to use otplus	e updated packages.			
5	Collecting package metadata (current_repodata.json):working done Note: you may need to restart the kernel to use updated packages. Solving environment:working done ## Package Plan ## environment location: C:\Users\rsnen\anaconda3 added / updated specs: - pydotplus						
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