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<b>8951 8952</b> 32	e_id	0.624 0.920 0.920	Male  Male  Female	No relevent experience  Has relevent experience  Has relevent experience  Has relevent experience  Has relevent experience	Full time course  no_enrollment  no_enrollment  no_enrollment	Graduate  Masters  Graduate  Graduate  Graduate	STEM STEM STEM Humanities
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######################################	0.762 0.920	of the state of th	21.0 21.0 21.0 21.0 21.0 21.0 21.0 21.0	with stra  2 2 6 6 3 2 2 3 3 1 2 2 3 3 1 2 2 3 3 1 2 2 3 3 1 2 2 4 4 5 6 6 6 7 8 8 6 8 6 8 9 6 8 9 6 8 9 8 9 8 9 8 9 8	size=0.2, randon  size=0.3, randon  size=0.4, randon  size=0.4, randon  size=0.5, randon  size=0.6, randon  size=0.7, randon  size=0.1, randon  size=0.2, randon  size=0.3, ra	poling are records in the test at the parameter of the test at the programment of the test and an institution of the test and at the parameter of the test and at the parameter of the test and the parameter of the test and the parameter of the test and the parameter of the pa	actify = y)  and, recall always are to describe a to decide and the classification module and all of these are the second and the classification module and all of these are the second and the classification and all of the second and the classification and all of the second and the classification and all of the second and the classification an
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	<pre>return models  # evaluate a given model using cross-validation  def evaluate_model(model, X, y):</pre>
	<pre>models = get_models() # evaluate the models and store results results, names = list(), list() for name, model in models.items():</pre>
	>5 0.811 (0.010) >6 0.815 (0.011) >7 0.816 (0.012) >8 0.819 (0.013) >9 0.829 (0.011) >10 0.831 (0.012) >11 0.834 (0.011) >12 0.838 (0.012) >13 0.836 (0.012) >14 0.838 (0.010) >15 0.838 (0.010) >16 0.842 (0.011) >17 0.843 (0.011) >18 0.850 (0.011) >19 0.851 (0.011) >20 0.852 (0.011) >20 0.852 (0.011) >21 0.852 (0.011) >22 0.853 (0.011) >23 0.853 (0.011) >24 0.853 (0.011) >25 0.854 (0.011) >27 0.854 (0.011) >28 0.854 (0.011) >29 0.854 (0.011) >29 0.854 (0.011) >29 0.854 (0.011) >29 0.854 (0.011) >29 0.854 (0.011) >29 0.854 (0.011) >29 0.854 (0.011)
	<pre>steps = [('pca', PCA(n_components=15)), ('m', KNeighborsClassifier(n_neighbors=5, p = 1, weights = "distance" model = Pipeline(steps=steps) model.fit(X_train, y_train) y_pred = model.predict(X_test) roc_auc_score(y_test, y_pred)  0.6740320139522297  All models perform similarly as they are based on the same platform, the knn. The main difference is in the optimization. Even the optimized models seem to not have much of an effect on the models accuracy on the test data, showing a better optimization algorithm is needed, potentially TPE algorithm.</pre> 7. Naive Bayes
In [141	<ol> <li>Train a model with GaussianNB, test it and then print the confusion matrix and classification report. Also, plot ROC curve and show the AUC of ROC, and the count of the number of misclassification.</li> <li>Train a model with CategoricalNB, test it and then print the confusion matrix and classification report. Also, plot ROC curve, and show the AUC of ROC and the count of the number of misclassification.</li> <li>from sklearn.naive_bayes import GaussianNB clf = GaussianNB() clf.fit(X_train, y_train) y_pred = clf.predict(X_test) print(confusion_matrix(y_test, y_pred))</li> </ol>
Tu [442	print(classification_report(y_test, y_pred))  [[1140 1102] [ 116 329]]
In [142	<pre>from sklearn.metrics import roc_curve fpr, tpr, thresholds = roc_curve(y_test, y_pred) def plot_roc_curve(fpr, tpr, label=None):     plt.plot(fpr, tpr, linewidth=2, label=label)     plt.plot([0, 1], [0, 1], 'k') # dashed diagonal  plot_roc_curve(fpr, tpr)  plt.xlabel("False positive rate") plt.ylabel("True positive rate (Recall)") plt.show()</pre>
	0.8 - (Recall) 0.6 - (Recall) 0.0 -
In [143 Out[143 In [144	<pre>from sklearn.metrics import roc_auc_score roc_auc_score(y_test, y_pred)  0.6239002094839079  from sklearn.naive_bayes import CategoricalNB clf = CategoricalNB() clf.fit(X_train, y_train) y_pred = clf.predict(X_test) print(confusion matrix(y test, y pred))</pre>
	print(classification_report(y_test, y_pred))  [[1902  340]
In [145	<pre>from sklearn.metrics import roc_curve fpr, tpr, thresholds = roc_curve(y_test, y_pred) def plot_roc_curve(fpr, tpr, label=None):     plt.plot(fpr, tpr, linewidth=2, label=label)     plt.plot([0, 1], [0, 1], 'k') # dashed diagonal  plot_roc_curve(fpr, tpr)  plt.xlabel("False positive rate") plt.ylabel("True positive rate (Recall)") plt.show()</pre>
	0.8 - 0.6 - 0.0 - 0.0 - 0.2 - 0.4 - 0.6 - 0.8 - 1.0 - 0.0 - 0.8 - 0.8 - 0.0 -
	from sklearn.metrics import roc_auc_score roc_auc_score(y_test, y_pred)  0.6960849562489351  Count of the number of misclassification  Gaussian = 1102 + 116  Categorical = 340 + 203  8. Support Vector Machine
In [39]:	<ol> <li>Build a support vector machine model using SVC. Use grid search to tune some parameters and then based on that show the best parameters found.</li> <li>Test the model and print the confusion matrix and classification report. Also, plot ROC curve and show the AUC of ROC, and the count of the number of misclassification.</li> </ol> from sklearn.svm import SVC import time
In [41]:	<pre>from sklearn.model_selection import KFold, StratifiedKFold from sklearn.model_selection import cross_val_score seed = 0 metric = 'accuracy' kFoldSplits = 5 np.random.seed(seed) best_score=1.0 def objective(space):  global best_score model = SVC(**space) kfold = StratifiedKFold(n_splits=kFoldSplits, random_state=seed, shuffle=True) # KFold is also an option score = 1-cross_val_score(model, X_train, y_train, cv=kfold, scoring=metric, verbose=False).mean() # Careful here (score). The objective function will be minimized, thus somme treatment on your score mig</pre>
In [42]:	<pre>if (score &lt; best_score):     best_score=score  return score  from hyperopt import hp, fmin, tpe, rand, STATUS_OK, Trials  space = {     'C': hp.choice('C', [0.1,1, 10, 100]),     'kernel': hp.choice('kernel',['sigmoid', 'poly', 'rbf']),     'gamma':hp.choice('gamma',[1,0.1,0.01,0.001])     } </pre>
In [43]:	<pre>n_iter_hopt = 10 trials = Trials() # Initialize an empty trials database for further saving/loading ran iteractions  start = time.time()  best = fmin(objective,</pre>
	elapsed_time_hopt = time.time() - start  100%
In [105 Out[105 In [106 In [107	<pre>model = svc(c = 1, gamma = 0.1, kernel = poly) model.fit(X_train,y_train)  SVC(C=1, gamma=0.1, kernel='poly')  y_pred = model.predict(X_test)</pre>
	[[1908
	<ol> <li>Decision Tree</li> <li>Build a decision tree model using sklearns DecisionTreeClassifier. Use the unbalanced training set, entropy as the criterion. Try with different max_depth (or use grid search). After building model, test it and print the confusion matrix and classification report. Also, plot ROC curve and show the AUC of ROC, and the count of the number of misclassification. Show the decision tree. (you can simply import tree from sklearn and call tree.plot_tree with your model and the call plt.show. At the beginning of this process, use plt.figure to change the figsize)</li> <li>Perform the same tasks as 9.1 with the balanced training set</li> </ol>
In [83]:	<pre>3. Discuss any difference and also discuss part of the tree of 9.2  from sklearn.tree import DecisionTreeClassifier def objective(space):  global best_score model = DecisionTreeClassifier(**space) kfold = StratifiedKFold(n_splits=kFoldSplits, random_state=seed, shuffle=True) # KFold is also an option score = 1-cross_val_score(model, X_unbalanced_train, y_unbalanced_train, cv=kfold, scoring=metric, verbox # Careful here (score). The objective function will be minimized, thus somme treatment on your score minimized if (score &lt; best_score):     best_score=score</pre>
In [117 In [85]:	<pre>space = {     'criterion': "entropy",         "max_depth": hp.choice("max_depth", range(1,51)) }  n_iter_hopt = 200 trials = Trials() # Initialize an empty trials database for further saving/loading ran iteractions</pre>
	<pre>start = time.time()  best = fmin(objective,</pre>
In [86]: In [87]: In [88]:	<pre>print("\nHyperopt search on unbalanced data took %.2f seconds for %d candidates. Accuracy reached: %.3f\nOption Hyperopt search on unbalanced data took 15.85 seconds for 200 candidates. Accuracy reached: 86.391 Optimal parameters found: {'max_depth': 2}  model = DecisionTreeClassifier(criterion = "entropy", max_depth = 2) model.fit(X_unbalanced_train, y_unbalanced_train) y_pred = model.predict(X_test)  from sklearn.metrics import classification_report, confusion_matrix print(confusion_matrix(y_test, y_pred))</pre>
	print(classification_report(y_test, y_pred))  [[2071 171] [ 205 240]]
In [90]:	<pre>from sklearn.metrics import roc_curve fpr, tpr, thresholds = roc_curve(y_test, y_pred) def plot_roc_curve(fpr, tpr, label=None):     plt.plot(fpr, tpr, linewidth=2, label=label)     plt.plot([0, 1], [0, 1], 'k') # dashed diagonal  plot_roc_curve(fpr, tpr)  plt.xlabel("False positive rate") plt.ylabel("True positive rate (Recall)") plt.show()</pre>
	1.0 - (Recall) 0.6 - (Recall) 0.4 - (Recall) 0.2 - (Recall) 0.4 - (Recall) 0.2 -
In [91]: Out[91]: In [118…	roc_auc_score(y_test, y_pred)  0.7315273281279756
In [119	<pre>global best_score   model = DecisionTreeClassifier(**space)   kfold = StratifiedKFold(n_splits=kFoldSplits, random_state=seed, shuffle=True) # KFold is also an option   score = 1-cross_val_score(model, X_train, y_train, cv=kfold, scoring=metric, verbose=False).mean()   # Careful here (score). The objective function will be minimized, thus somme treatment on your score mig  if (score &lt; best_score):     best_score=score  return score  n iter hopt = 200</pre>
	<pre>trials = Trials() # Initialize an empty trials database for further saving/loading ran iteractions  start = time.time()  best = fmin(objective,</pre>
In [120 In [125	print("\nHyperopt search on balanced data took %.2f seconds for %d candidates. Accuracy reached: %.3f\nOptimes  Hyperopt search on balanced data took 36.11 seconds for 200 candidates. Accuracy reached: 86.391  Optimal parameters found: {'max_depth': 33}  model = DecisionTreeClassifier(criterion = "entropy", max_depth = 34)  model.fit(X_train, y_train)  y_pred = model.predict(X_test)
In [126	<pre>from sklearn.metrics import classification_report, confusion_matrix print(confusion_matrix(y_test, y_pred)) print(classification_report(y_test, y_pred))  [[1890</pre>
	0.0 0.88 0.84 0.86 2242 1.0 0.35 0.42 0.38 445 accuracy 0.77 2687 macro avg 0.61 0.63 0.62 2687 weighted avg 0.79 0.77 0.78 2687
In [127	0.0
	0.0 0.88 0.84 0.86 2242 1.0 0.35 0.42 0.38 445  accuracy 0.77 2687 macro avg 0.61 0.63 0.62 2687 weighted avg 0.79 0.77 0.78 2687  Misclassification = 352 + 258  from sklearn.metrics import roc_curve fpr, tpr, thresholds = roc_curve(y_test, y_pred) def plot_roc_curve(fpr, tpr, label=None):     plt.plot(fpr, tpr, linewidth=2, label=label)     plt.plot([0, 1], [0, 1], 'k') # dashed diagonal  plot_roc_curve(fpr, tpr)
In [128 Out[128	accuracy macro avg 0.61 0.63 0.62 242 macro avg 0.61 0.63 0.62 2887 weighted avg 0.79 0.77 0.78 2887  Misclassification = 352 + 258   from sklearn.metrics import roc curve fpr, tpr, thresholds = roc_curve(y_test, y_pred) def plot_roc_curve(fpr, tpr, linewidthey, label=label)     plt.plot(fpr, tpr, linewidthey, label=label)     plt.plot(fpr, tpr, linewidthey, label=label)     plt.plot(fpr, tpr, linewidthey, label=label)     plt.ylabel("Talae positive rate")     plt.ylabel("Talae positive rate (Recall)")     plt.show()  from sklearn.metrics import roc_auc_score roc_auc_score(y_test, y_pred)  0.6316110214595716  This difference in acuracy could be due to the imbalance present in the testing set not present in the training set as it was ballanced. This
In [128 Out[128	accuracy 1.0 0.88 0.84 0.86 2747 1.0 0.23 0.42 0.38 445  accuracy 1.0 0.23 0.42 0.38 445  3.77 1887  metaphoracy 1.61 0.81 0.82 2987  Misclassification = 352+258  from ski-earn.metrics import roc_curve ghr, bpr, bbrnshalds = roc_corvety_total_y_sred) def plor_roc_curve(fpr, tpr, label=None): ght.piot(fpr, lpr, threshalds) = roc_corvety_total_y_sred) def plor_roc_curve(fpr, tpr, label=None): ght.piot(fpr, lpr, threshalds) = roc_corvety_total_y_sred) def plor_roc_curve(fpr, tpr, label=None): ght.piot(fpr, lpr, threshalds) = roc_corvety_total_y_sred) plor_roc_curve(fpr, tpr, label=None): ght.piot(fpr, lpr, threshalds) = roc_corvety_total_y_sred) def plor_roc_curve(fpr, tpr, label=None): ght.piot(fpr, lpr, threshalds) plor_poc_curve(fpr, tpr, lpr, label=None): ght.piot(fpr, lpr, threshalds) plor_poc_curve(fpr, tpr, label=None): ght.piot(fpr, lpr, label=None): ght.p
In [128 Out[128	accuracy macro and 0.64 0.86 2042  accuracy macro and 0.61 0.65 0.62 2897  watched awd 0.79 0.77 0.78 0.687  Misclassification = 352 + 258  from **Khorm.metrics import ring curve (picture) (pictur
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