```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
import warnings
warnings.simplefilter(action='ignore', category=FutureWarning)
y = pd.read csv('/content/actual.csv')
X train = pd.read csv('/content/data set ALL AML train.csv')
X test = pd.read csv('/content/data set ALL AML independent.csv')
# Remove "call" columns from training and testing data
train to keep = [col for col in X train.columns if "call" not in col]
test to keep = [col for col in X test.columns if "call" not in col]
X train tr = X train[train to keep]
X test tr = X test[test to keep]
'27', '28', '29', '30', '31', '32', '33', '34', '35',
      '26',
'36', '37', '38']
X train tr = X train tr.reindex(columns=train columns titles)
test columns titles = ['Gene Description', 'Gene Accession
Number', '39', '40', '41', '42', '43', '44', '45', '46', '47', '48', '49', '50', '51', '52', '53', '54', '55', '56',
'70', '71', '72']
X test tr = X test tr.reindex(columns=test columns titles)
X train = X train tr.T
X \text{ test} = X \text{ test tr.T}
X train.head()
{"type":"dataframe", "variable name": "X train"}
X test.head()
{"type": "dataframe", "variable name": "X test"}
# Clean up the column names for training and testing data
X train.columns = X train.iloc[1]
X train = X train.drop(["Gene Description", "Gene Accession
Number"]).apply(pd.to numeric)
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# Clean up the column names for Testing data
X test.columns = X test.iloc[1]
X test = X test.drop(["Gene Description", "Gene Accession
Number"]).apply(pd.to numeric)
X train.head()
{"type": "dataframe", "variable name": "X train"}
from sklearn.preprocessing import LabelEncoder
le = LabelEncoder()
y["cancer"] = le.fit transform(y["cancer"])
У
{"summary":"{\n \"name\": \"y\",\n \"rows\": 72,\n \"fields\": [\n \]}
{\n \"column\": \"patient\",\n \"properties\": {\n \"dtype\": \"number\",\n \"std\": 20,\n \"min\": 1,\n
\"max\": 72,\n \"num_unique_values\": 72,\n [\n 5 \n 63 \n 19\n]
                                                             \"samples\":
             5,\n
                                                         ],\n
[\n
                          63.\n
                                            19\n
\"semantic_type\": \"\",\n \"description\": \"\"\n
            {\n \"column\": \"cancer\",\n \"properties\":
{\n \"dtype\": \"number\",\n \"std\": 0,\n
\"min\": 0,\n \"max\": 1,\n \"num_unique_val
\"samples\": [\n 1,\n 0\n ],\n
\"semantic_type\": \"\",\n \"description\": \"\"\n
                                           \"num unique values\": 2,\n
                                                                  }\
     }\n ]\n}","type":"dataframe","variable name":"y"}
X train["patient"] = X train.index.values
X test["patient"] = X test.index.values
X train = X train.astype("int32")
X test = X test.astype("int32")
X train
{"type": "dataframe", "variable name": "X train"}
{"summary":"{\n \"name\": \"y\",\n \"rows\": 72,\n \"fields\": [\n
{\n \"column\": \"patient\",\n \"properties\": {\n
\"dtype\": \"number\",\n \"std\": 20,\n \"min\": 1,\n
             5,\n
                       \"num unique_values\": 72,\n
                                                             \"samples\":
\"max\": 72,\n
[\n
                          63,\n
                                            19\n
                                                         ],\n
\"semantic_type\": \"\",\n \"description\": \"\"\n
n },\n {\n \"column\": \"cancer\",\n \"properties\":
           \"dtype\": \"number\",\n \"std\": 0,\n
{\n
```

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\"min\": 0,\n \"max\": 1,\n \"samples\": [\n 1,\n
                                          \"num unique values\": 2,\n
                                         0\n
                                                   ],\n
\"semantic_type\": \"\",\n \"description\": \"\"\n
                                                                }\
     }\n ]\n}","type":"dataframe","variable_name":"y"}
X train = pd.merge(X train, y, on="patient")
X test = pd.merge(X test, y, on="patient")
X train["cancer"].value counts()
cancer
     27
1
     11
Name: count, dtype: int64
X_test["cancer"].value_counts()
cancer
     20
     14
1
Name: count, dtype: int64
import random
upsampled_data = random.sample(X_train.query("cancer == 1")
["patient"].index.to_list(), k=8, )
train data upsampled = pd.concat([X train,
X train.iloc[upsampled data, :]])
X train = train data upsampled.drop(columns=["patient", "cancer"])
y train = train data upsampled["cancer"]
y_test = X test["cancer"]
X test = X test.drop(columns=["patient", "cancer"])
from sklearn.preprocessing import StandardScaler
sc = StandardScaler()
X_train_scaled = sc.fit_transform(X_train)
X test scaled = sc.transform(X test)
from sklearn.model selection import cross val score
from sklearn.ensemble import RandomForestClassifier
from sklearn.model selection import GridSearchCV
parameters = {"bootstrap": [False, True],
             "n_estimators": [60, 70, 80, 90, 100],
             "max_features": [0.6, 0.65, 0.7, 0.75, 0.8],
             "min samples leaf": [8, 10, 12, 14],
             "min samples split": [3, 5, 7]
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}
model = RandomForestClassifier()
cv = cross val score(estimator=model,
X=X train scaled,
y=y_train,
cv=3,
scoring="accuracy")
rf GS = GridSearchCV(model, parameters,cv=3, scoring="accuracy")
rf GS.fit(X train scaled, y train)
best rf = rf GS.best estimator
                                           Traceback (most recent call
NameError
last)
<ipython-input-35-c4f7a56edccc> in <cell line: 23>()
     21 rf GS = GridSearchCV(model, parameters,cv=3,
scoring="accuracy")
     22 rf GS.fit(X train scaled, y train)
---> 23 best rf = rf search.best estimator
NameError: name 'rf search' is not defined
best rf
RandomForestClassifier(bootstrap=False, max features=0.6,
min samples leaf=14,
                       min_samples_split=7, n_estimators=70)
import sklearn.metrics as metrics
rf prediction = best rf.predict(X test scaled)
accuracy = metrics.accuracy_score(y_test, rf_prediction)
accuracy
0.9117647058823529
from sklearn.neighbors import KNeighborsClassifier
parameters = {
    "n neighbors": [i for i in range(1, 30, 5)],
    "weights": ["uniform", "distance"],
    "algorithm": ["kd tree"],
    "leaf size": [1, \overline{10}, 20, 30],
    "p": [1, 2]
}
model = KNeighborsClassifier()
```

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cv = cross val score(estimator=model,
X=X train scaled,
 y=y_train,
 cv=3,
 scoring="accuracy")
knn GS = GridSearchCV(KNeighborsClassifier(), parameters, n jobs=-1,
verbose=1, scoring="accuracy")
knn GS.fit(X train scaled, y train)
best knn = knn GS.best estimator
Fitting 5 folds for each of 96 candidates, totalling 480 fits
best knn
KNeighborsClassifier(algorithm='kd tree', leaf size=1, n neighbors=6,
p=1,
                     weights='distance')
knn prediction = best knn.predict(X test scaled)
accuracy = metrics.accuracy score(y test, knn prediction)
accuracy
0.7647058823529411
from sklearn.linear model import LogisticRegression
parameters = {"C": [1e-03, 1e-2, 1e-1, 1, 10],
      "penalty": ["l1", "l2"]}
model = LogisticRegression()
cv = cross_val_score(estimator=model,
 X=X train scaled,
y=y_train,
 cv=3,
 scoring="accuracy")
log GS = GridSearchCV(model, parameters, cv=3, scoring="accuracy")
log GS.fit(X train scaled, y train)
best lr = log GS.best estimator
/usr/local/lib/python3.10/dist-packages/sklearn/model selection/
validation.py:378: FitFailedWarning:
15 fits failed out of a total of 30.
The score on these train-test partitions for these parameters will be
set to nan.
If these failures are not expected, you can try to debug them by
setting error score='raise'.
```

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Below are more details about the failures:
15 fits failed with the following error:
Traceback (most recent call last):
  File
"/usr/local/lib/python3.10/dist-packages/sklearn/model selection/ vali
dation.py", line 686, in _fit_and_score
    estimator.fit(X train, y train, **fit params)
  File
"/usr/local/lib/python3.10/dist-packages/sklearn/linear_model/_logisti
c.py", line 1162, in fit
    solver = _check_solver(self.solver, self.penalty, self.dual)
"/usr/local/lib/python3.10/dist-packages/sklearn/linear_model/_logisti
c.py", line 54, in check solver
    raise ValueError(
ValueError: Solver lbfgs supports only 'l2' or 'none' penalties, got
ll penalty.
 warnings.warn(some_fits_failed_message, FitFailedWarning)
/usr/local/lib/python3.10/dist-packages/sklearn/model selection/ searc
h.py:952: UserWarning: One or more of the test scores are non-finite:
        nan 0.95833333
                              nan 0.95833333
                                              nan 0.95833333
        nan 0.95833333
                              nan 0.9375 l
 warnings.warn(
best lr = log GS.best estimator
lr prediction = best lr.predict(X test scaled)
accuracy = metrics.accuracy score(y test, lr prediction)
accuracy
0.8235294117647058
from sklearn.tree import DecisionTreeClassifier
parameters = {'max leaf nodes': list(range(2, 100)),
'min samples split': [2, 3, 4, 5, 6], 'max_depth':[3,4,5,6,7,8]}
model = DecisionTreeClassifier()
cv = cross val score(estimator=model,
X=X train scaled,
y=y_train,
cv=3,
 scoring="accuracy")
dtc GS = GridSearchCV(model, parameters, verbose=1, cv=3,
scoring="accuracy")
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dtc GS.fit(X train scaled, y train)
best dtc = dtc GS.best estimator
Fitting 3 folds for each of 2940 candidates, totalling 8820 fits
NameError
                                             Traceback (most recent call
last)
<ipython-input-61-80555a0ae8bb> in <cell line: 15>()
     13 dtc GS = GridSearchCV(model, parameters, verbose=1, cv=3,
scoring="accuracy")
     14 dtc GS.fit(X train scaled, y train)
---> 15 best dtc = dst search.best estimator
NameError: name 'dst_search' is not defined
best dtc
DecisionTreeClassifier(max depth=3, max leaf nodes=3,
min samples split=4)
dtc prediction = best dtc.predict(X test scaled)
accuracy = metrics.accuracy score(y test, dtc prediction)
accuracy
0.9117647058823529
rf acc = metrics.accuracy score(y test,
best rf.predict(X test scaled))
knn acc = metrics.accuracy score(y test,
best knn.predict(X test scaled))
lr acc = metrics.accuracy score(y test,
best_lr.predict(X_test scaled))
dtc acc = metrics.accuracy score(y test,
best dtc.predict(X test scaled))
model acc = pd.DataFrame({
    'Model': ['KNN', 'Logistic Regression', 'Decision Tree',
               'Random Forest'],
    'accuracy score': [knn acc, lr acc, dtc acc,
               rf acc]})
model acc.sort values(by='accuracy score', ascending=False)
{"summary":"{\n \"name\": \"model_acc\",\n \"rows\": 4,\n
\"fields\": [\n {\n \"column\": \"Model\",\n \"properties\": {\n \"dtype\": \"string\",\n
\"num_unique_values\": 4,\n \"samples\": [\n
Forest\",\n \"KNN\",\n \"Decision Tree\"\
n ],\n \"semantic_type\": \"\",\n
                                                                  \"Random
```

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\"description\": \"\"\n
                                                   \"column\":
                                  },\n {\n
                           }\n
\"accuracy score\",\n \"properties\": {\n
                                                   \"dtype\":
\"number\",\n \"std\": 0.07204381596421114,\n
                                                         \"min\":
0.7647058823529411,\n\\"max\": 0.9117647058823529,\n
\"num_unique_values\": 3,\n \"samples\": [\n 0.9117647058823529,\n 0.8235294117647058,\n
0.7647058823529411\n
                           ],\n \"semantic type\": \"\",\n
\"description\": \"\"\n }\n ]\n}","type":"dataframe"}
mean_acc = np.mean([rf_acc, knn_acc, lr_acc, dtc_acc])
weight rf = rf acc / mean acc
weight knn = knn acc / mean acc
weight lr = lr acc / mean acc
weight_dt = dtc_acc / mean_acc
model weights = pd.DataFrame({
    'Model': ['KNN', 'Logistic Regression', 'Decision Tree',
             'Random Forest'],
    'weight': [weight knn, weight lr, weight dt,
             weight rf]})
model weights.sort values(by='weight', ascending=False)
{"summary":"{\n \"name\": \"model weights\",\n \"rows\": 4,\n
\"fields\": [\n \\"column\\": \\"Model\\",\n
\"properties\": {\n \"dtype\": \"string\",\n
\"num_unique_values\": 4,\n \"samples\": [\n
Forest\",\n \"KNN\",\n \"Decision Tree\"\
                                                           \"Random
Forest\",\n \"KNN\",\n \"Decis
n ],\n \"semantic_type\": \"\",\n
\"column\":
\"weight\",\n \"properties\": {\n
                                            \"dtype\": \"number\",\n
\"std\": 0.08446516354424746,\n\\"min\": 0.896551724137931,\n\\"max\": 1.0689655172413792,\n\\"num_unique_values\": 3,\n
\"semantic_type\": \"\",\n \"description\": \"\"\n }\
    }\n ]\n}","type":"dataframe"}
from sklearn.ensemble import VotingClassifier
ensemble = VotingClassifier(estimators=[("rf", best rf), ("knn",
best knn), ("lr", best lr), ("dt", best dtc)],
                                      voting="soft",
weights=[weight rf, weight knn, weight_lr, weight_dt])
ensemble.fit(X train scaled, y train)
VotingClassifier(estimators=[('rf',
                             RandomForestClassifier(bootstrap=False,
                                                   max features=0.6,
```

```
min samples leaf=14,
min samples split=7,
n estimators=70)),
                             ('knn',
KNeighborsClassifier(algorithm='kd_tree',
                                                    leaf size=1,
n_neighbors=6,
                                                    p=1,
weights='distance')),
                             ('lr', LogisticRegression(C=0.001)),
                             ('dt',
                              DecisionTreeClassifier(max depth=3,
                                                      max leaf nodes=3,
min_samples_split=4))],
                 voting='soft',
                 weights=[1.0689655172413792, 0.896551724137931,
                          0.9655172413793104, 1.0689655172413792])
ens prediction = ensemble.predict(X test scaled)
acc_score = metrics.accuracy_score(y_test, ens_prediction)
acc_score
0.9411764705882353
```