

Goals

- Laid out a rough outline of how SHAP would be computed, I thought I would give SHAP methods a try
- Earlier methods work and prove that the model is unpickled and can be used
- Be able to iterate through each trained model to it's respective CV dataset, create shap values, generate shap plots
- Be able to store each CV shap values for each model and store in csv file as a DataFrame

```
LR_shap_all_CVs.csv ==>
                        LR_0 --> CV0
                        LR_1 --> CV1
                        LR_2 --> CV2
```

Things to do:

- Still need to figure out saving results into a file (pickle.dump()), create and save into designated folder
- Figure out how to work TreeExplainer, expected_value function
- Find file with the feature names for corresponding dataset to load into program under 'Load Metadata" section
- Figure out how to display other shap plots such as waterfall, force plot, etc

Notes

- Most of the program is hardcoded to specifically load one of the trained models after running STREAMLINE **resolved**
- Was able to prove that the model can be unpickled and used for .predict() and .predictproba() **resolved**
- Was able to use model to create SHAP explainers, calculate shap_values for CV0 testing dataset, and display plots **resolved**
- However, still need to refine the SHAP methods as there were some issues for Decision Tree Classifier **resolved**
- Was able to display Decision Tree prediction using TreeExplainer or even Explainer....I might be doing something wrong **resolved**
- XGBOOST MODEL IS COMPATIBLE WITH ALL OF THE LISTED SHAP PLOTS **resolved**
- RF MODEL NEEDED IT'S OWN IF-STATEMENT FOR NOW BUT WILL CONDENSE FOR CLARITY ADN EFFICIENCY **resolved**
- STILL NEED TO WORK ON LIGHTGBM, CATBOOST **resolved**
- GO BACK TO FIX DECISION TREE **resolved**

Fix

- Go back to double check shap plot compatibility for global and local importance for linear models **resolved**
- Work through the DecisionTreeClassifier and compare to other codes out there (if possible) **resolved**
- Currently unsure if creating dataframe for each model's shap_values shuold be done in compute_shap_values() or within the nested for-loop in testing cell
- Feature names when displaying shap plots

Updates (refer to 'Next Steps' for more updates)

7/29/22

- ALL given SHAP plots seems to work for NB() when not in a defined function block and if-statement **resolved**
- Bar, scatter, waterfall, and beeswarm plots don't work for LR(), other plots work fine on LinearExplainer() and shap_values = explainer.shap_values(data)

8/02/22

- Plots and shap_values for each trained model in each CV work
- Will focus on section called '**Next Steps**'
 - refer to bottom of Notebook for more details
 - Currently unsure if creating dataframe for each model's shap_values shuold be done in compute_shap_values() or within the nested for-loop in testing cell

```
In [1]: # required packages & models
import os
import sys
import glob
import pickle
import warnings
warnings.filterwarnings('ignore')
import csv
import sklearn
import shap
import numpy as np
import numpy.typing as npt
```

```
import pandas as pd
import scipy as sp
import matplotlib.pyplot as plt
from matplotlib.backends.backend_pdf import PdfPages
from fpdf import FPDF
import collections
from termcolor import colored as cl #text customization

# Model packages
import xgboost
import lightgbm as lgb
from sklearn import *
from sklearn.naive_bayes import GaussianNB
from sklearn.linear_model import LogisticRegression
from sklearn.tree._classes import DecisionTreeClassifier
from sklearn.ensemble import RandomForestClassifier
import xgboost as xgb
import lightgbm as lgb
import catboost as cgb
from sklearn import tree
from shap.plots import waterfall

#import metrics
from sklearn.model_selection import train_test_split
from sklearn.metrics import classification_report, accuracy_score

# Jupyter Notebook Hack: This code ensures that the results of multiple commands within a given cell are all displayed
from IPython.core.interactiveshell import InteractiveShell
InteractiveShell.ast_node_interactivity = "all"

shap.initjs() # load JS visualization code to notebook. SHAP plots won't be displayed without this
```



Run Parameters

```
In [2]: dataset_path = "/Users/jessicakim/Desktop/STREAMLINE/DemoData"
experiment_path = "/Users/jessicakim/Desktop/STREAMLINE/DemoData/Output/hcc_demo"
targetDataName = 'None'

# hardcoded pathways for CVDataset0
# train_file_path = '/hcc-data_example/CVDatasets/'
# test_file_path = '/hcc-data_example/CVDatasets/'
```

Check for Analyzed Datasets and Remove Unecessary Files

```
In [3]: datasets = os.listdir(experiment_path)
experiment_name = experiment_path.split('/')[-1] #Name of experiment folder

datasets.remove('metadata.csv')
datasets.remove('metadata.pickle')
datasets.remove('algInfo.pickle')

try:
    datasets.remove('jobsCompleted')
except:
    pass
try:
    datasets.remove('UsefulNotebooks')
except:
    pass
try:
    datasets.remove('logs')
    datasets.remove('jobs')
except:
    pass
try:
    datasets.remove('DatasetComparisons') #If it has been run previously (overwrite)
except:
    pass
try:
    datasets.remove('KeyFileCopy') #If it has been run previously (overwrite)
except:
    pass
try:
    datasets.remove('.DS_Store') #If it has been run previously (overwrite)
except:
    pass
try:
    datasets.remove(experiment_name+'_ML_Pipeline_Report.pdf') #If it has been run previously (overwrite)
except:
    pass

datasets = sorted(datasets) #ensures consistent ordering of datasets
print("Analyzed Datasets: "+str(datasets))

Analyzed Datasets: ['hcc-data_example', 'hcc-data_example_no_covariates']
```

Load Metadata and Other Necessary Variables

```
In [4]: jupyterRun = 'True'
# Loading necessary variables specified earlier in the pipeline from metadatafor dataPrep()
file = open(experiment_path + '/' + "metadata.pickle", 'rb')
metadata = pickle.load(file)
# file.close()
# print(metadata)

class_label = metadata['Class Label']
instance_label = metadata['Instance Label']
cv_partitions = int(metadata['CV Partitions'])

# # # unpickle and load in feature_names found in 'categorical_variables.pickle'
# feature_names_file = experiment_path + '/hcc-data_example/exploratory/categorical_variables.pickle'
# file = open(feature_names_file , 'rb')
# feature_names= pickle.load(file)
# file.close()
# print('Checking for feature names...\n',feature_names)

alg_file = open(experiment_path + '/' + "/algInfo.pickle", 'rb')
algInfo = pickle.load(alg_file)
alg_file.close()
algorithms = []

abbrev = {}
for key in algInfo: # pickling specific model while also checking for corresponding algInfo
    if algInfo[key][0]: # If that algorithm was used
        algorithms.append(key)
        abbrev[key] = (algInfo[key][1])

print('\nChecking for algorithms used in STREAMLINE...\n',algorithms)
print('\nChecking for abbrev for algorithms used in STREAMLINE...\n', abbrev)
```

Checking for algorithms used in STREAMLINE...
['Naive Bayes', 'Logistic Regression', 'Decision Tree', 'Random Forest', 'Extreme Gradient Boosting']

Checking for abbrev for algorithms used in STREAMLINE...
{'Naive Bayes': 'NB', 'Logistic Regression': 'LR', 'Decision Tree': 'DT', 'Random Forest': 'RF', 'Extreme Gradient Bo
osting': 'XGB'}

Get Feature Names From Target Dataset

```
In [5]: # user can choose which csv dataset file to use if more than one was analyzed
target_dataset = '/hcc-data_example.csv' # default is 'None'
orig_dataset = dataset_path + '/' + target_dataset
# print(orig_dataset)

# feature_names = pd.read_csv(orig_dataset)
# if instance_label != 'None':
#     feature_names = feature_names.drop(instance_label,axis=1)
#     feature_names = feature_names.drop(class_label, axis= 1).columns
# print(feature_names)
```

dataPrep(): Loading Target CV Training & Testing Sets

```
In [6]: def dataPrep(train_file_path,instance_label,class_label, test_file_path):

    '''Loads target cv training dataset, separates class from features and removes instance labels'''

    # store all used features from training and testing sets which will be later used for shapley valeues & plots
    train_feat = {}
    test_feat = {}

    train = pd.read_csv(train_file_path)
    if instance_label != 'None':
        train = train.drop(instance_label,axis=1)

    # populate train_feat[] with feature names from given train dataset
    train_feat = train.drop(class_label, axis = 1).columns.tolist()
    print(train_feat)

    trainX = pd.DataFrame(train.drop(class_label,axis=1).values)
    trainY = pd.DataFrame(train[class_label].values)
    del train #memory cleanup

    test = pd.read_csv(test_file_path)
    if instance_label != 'None':
        test = test.drop(instance_label,axis=1)
```

```
# populate train_feat[] with feature names from given test dataset
test_feat = test.drop(class_label, axis = 1).columns.tolist()
print(test_feat)

testX = pd.DataFrame(test.drop(class_label,axis=1).values)
testY = pd.DataFrame(test[class_label].values)
del test #memory cleanup

return trainX, trainY, testX, testY, train_feat, test_feat
```

SHAP: get_explainer()

- will check if explainer is one of the available ML in STREAMLINE
- if algorithm name matches ['list model names'], create explainers
- return explainer based on given model from parameter

Types of SHAP Explainers

.Explainer()

- Uses Shapley values to explain any machine learning model or python function.
- This is the primary explainer interface for the SHAP library
- It takes any combination of a model and masker and returns a callable subclass object that implements the particular estimation algorithm that was chosen.

.TreeExplainer()

- Uses Tree SHAP algorithms to explain the output of ensemble tree models.
- Tree SHAP is a fast and exact method to estimate SHAP values for tree models and ensembles of trees, under several different possible assumptions about feature dependence.
- It depends on fast C++implementations either inside an external model package or in the local compiled C extention.

.LinearExplainer()

- Computes SHAP values for a linear model, optionally accounting for inter-feature correlations.
- This computes the SHAP values for a linear model and can account for the correlations among the input features.
- Assuming features are independent leads to interventional SHAP values which for a linear model are $coef[i] * (x[i] - X.mean(0)[i])$ for the i th feature.
- If instead we account for correlations then we prevent any problems arising from colinearity and share credit among correlated features.
- Accounting for correlations can be computationally challenging, but LinearExplainer uses sampling to estimate a transform that can then be applied to explain any prediction of the model.

```
In [7]: def get_explainer(model, abbrev, trainX):

    '''Pass loaded model and abbrev to match appropriate SHAP explainer'''

    '''Must always use training dataset as background data in order to
        evaluate SHAP values for either testing (usually) or training set'''

    explainer = None
    trained_model = model

    if abbrev in ["NB"]:
        explainer = shap.Explainer(trained_model.predict, trainX)

        # dont use model.predict for Linear Explainer (only for Explainer)
        # ^^^ You get a class method error when creating shap plots and values
    if abbrev in ["LR"]:
        explainer = shap.LinearExplainer(trained_model, trainX)

    if abbrev in ['DT', 'RF', "XGB", "LGB","CGB"]:
        explainer = shap.TreeExplainer(trained_model)

    return explainer
```

SHAP: compute_shapValues()

NOTES

- Parameter 'X' in this context refers to whatever training or testing dataset that was passed in from the whole run from below
- Mentioned earlier, default run uses training dataset as background data and creates shap values using testing data
- The same follows for feature_names --> either train_feat or test_feat (default) will be passed

```
In [8]: def compute_shapValues(model, abbrev, explainer, X) -> pd.DataFrame :  
  
    '''This method will calculate shapley values and store these as a Pandas DataFrame for conversion to csv file  
        This includes creating expected_values and shap_values --> returns shap_values (will be called by shap_summary)  
    '''  
  
    max_evals = max(500, (2 * len(X)) + 1)    # optional: declares number of permutations for shap.Explainer()  
    shap_values = None  
  
    if abbrev in ["NB"]:  
        shap_values= explainer(X)    # permutation object cannot use .expected_value function like LR  
        print(shap_values)  
  
  
    if abbrev in ["LR"]:  
        shap_values = explainer.shap_values(X)  
        print(shap_values)  
  
  
    # i think shap_values() only works for TreeExplainer and LinearExplainer...Explainer for NB is considered a  
    # permutation object  
    if abbrev in ['DT', 'RF', "XGB", "LGB","CGB"]:  
        shap_values = explainer.shap_values(X, approximate=False, check_additivity=False)  
        print(shap_values)  
  
  
    return shap_values
```

SHAP: shap_summary()

Plot Types for SHAP v0.41.0

Waterfall

- Plots an explantion of a single prediction as a waterfall plot

Summary (type: violin & bar)

- Summary plots of SHAP values across a whole dataset

Dependence

- Plots the value of the feature on the x-axis and the SHAP value of the same feature on the y-axis
- This shows how the model depends on the given feature, and is like a richer extenstion of the classical parital dependence plots.
- Vertical dispersion of the data points represents interaction effects.
- Grey ticks along the y-axis are data points where the feature's value was NaN.

Force

- Visualize cumulative SHAP values with an additive force layout.

Beeswarm

- Summary plots of SHAP values across a whole dataset
- Designed to display an information-dense summary of how the top features in a dataset impact the model's output.

```
In [9]: def shap_summary(abbrev, feature_names,shap_values, explainer, X, cvCount):  
    '''Retrieve shap_values from previous method;  
        this method will return and display different types of shap plots  
  
        Figures for each model CV is saved as a png which will be merged to a  
        final summary report for each model  
    '''  
  
    save_path = experiment_path + '/hcc-data_example/model_evaluation/shap_values/testResults/shapFigures/'  
    # checks algorithm in given list to execute shap summaries  
    if abbrev in ["NB"]:  
  
        print('Summary Plot for SHAP Values in Class 0 & 1 in Test Set: \n')  
        shap.summary_plot(shap_values, X, feature_names, plot_type='violin', show=False)  
        # print('SHAP Bar Plot for Summary Plot for SHAP Values in Class 0 & 1 in Test Set:\n')  
#        shap.plots.bar(shap_values.values) # doesnt work but should for this...attribute error  
  
        print('SHAP Beeswarm Plot for Top 5 SHAP Values in Class 0 & 1 in Test Set: \n')  
        shap.plots.beeswarm(shap_values, max_display=5, show=False)    #max_display allows user to choose # of features
```



```

        # print('Waterfall Plot for SHAP Values in Class 0 in Test Set: \n')
#         shap.waterfall_plot(shap_values, max_display=5, show=True) # should work for this model


# #         # scatter, bar, waterfall, beeswarm plots should work for this model
# #         # waterfall plot also doesnt work...i get "AttributeError: 'numpy.ndarray' object has no attribute 'base_v
# #         Bar plot should work for this model if using .Explainer() and shap_values = explainer(data)-->
# #         not explainer.shap_values
elif abbrev in ["LR", 'XGB']:

    expected_value = explainer.expected_value
    print('Expected value for {}: {}'.format(abbrev, expected_value))

    print('Summary Plot for SHAP Values in Test Set: \n')
    shap.summary_plot(shap_values, X, feature_names, plot_type='violin', show=False)
    plt.savefig(save_path+abbrev+'_'+str(cvCount)+'shapSummaryPlot.png', bbox_inches='tight')
    plt.close()

    print('SHAP Bar Plot for SHAP Values Test Set: \n')
    shap.summary_plot(shap_values, X, feature_names, plot_type="bar", show=False)
    plt.savefig(save_path+abbrev+'_'+str(cvCount)+'shapSummaryBarPlot.png', bbox_inches='tight')
    plt.close()

    print('SHAP Decision Plot for SHAP Values in Test Set: \n')
    shap.decision_plot(expected_value, shap_values, feature_names, show=False)
    plt.savefig(save_path+abbrev+'_'+str(cvCount)+'shapDecisionPlot.png', bbox_inches='tight')
    plt.close()

    print('SHAP Decision Plot for Single-Prediction in Test Set: \n')
    shap.decision_plot(expected_value, shap_values[54], feature_names, show=False)
    plt.savefig(save_path+abbrev+'_'+str(cvCount)+'shapDecisionPlot_singlePredict.png', bbox_inches='tight')
    plt.close()

    # waterfall plot works for DT() if it uses .Explainer() and shap_vales = explainer(data)
    # instead of using TreeExplainer but other plots listed here work
elif abbrev in ['DT', 'RF', 'LGB', 'CGB']:
    expected_value = explainer.expected_value
    print('Expected value for {}: {}'.format(abbrev, expected_value))

    print('Bar Summary Plot for SHAP Values in Class 0 & 1 in Test Set: \n')
    #         #tree.tree_plot(testX) ---> helps display Decision Tree
    shap.summary_plot(shap_values, X, feature_names, plot_type='bar', class_names=['0', '1'], show=False)

    print('\nDecision Plot for SHAP Values from Class 0 in Test Set: \n')
    shap.decision_plot(expected_value[0], shap_values[0], feature_names=feature_names, show=False)

    print('\nDecision Plot for SHAP Values from Class 1 in Test Set: \n')
    shap.decision_plot(expected_value[1], shap_values[1], feature_names=feature_names, show=False)
```

This cell below might be to create summary reports

In []:

```
In [10]: # def run_force_plot(model, abbrev, explainer, shap_values, trainX, testX, run = True):
#         if abbrev in ['NB']:
#             print('\nForce Plot for {} SHAP Values from Class 0 in Test Set: \n'.format(abbrev))
#             shap.force_plot(shap_values[0], testX.iloc[0], feature_names=feature_names, show=True)

#             print('\nForce Plot for {} SHAP Values from Class 0 in Test Set: \n'.format(abbrev))
#             shap.force_plot(shap_values[1], testX.iloc[1], feature_names=feature_names, show=True)

#         elif abbrev in ['LR', 'XGB']:
#             print('\nChecking if shap plots are returned and consistent...\n')
#             summary = shap_summary(algorithms, shap_values, explainer, trainX, testX) # retrieve shap summary plots

#             print('\nForce Plot for SHAP Values in Whole Test Set: \n')
#             shap.force_plot(explainer.expected_value, shap_values, testX)

#         else:
#             print('\nForce Plot for {} SHAP Values from Class 0 in Test Set: \n'.format(abbrev))
#             shap.force_plot(explainer.expected_value[0], shap_values[0], feature_names=feature_names)

#             print('\nForce Plot for {} SHAP Values from Class 1 in Test Set: \n'.format(abbrev))
#             shap.force_plot(explainer.expected_value[1], shap_values[1], feature_names=feature_names)
```

^^^ fix later ... may want to keep this to create force plots and save results

```
In [11]: def shap_feature_ranking(abbrev, shap_values, X, feature_names): # 'X' and 'feature_names' argument is whichever test
```

```
'''Calculate the average of the absolute SHAP values for each feature and use it to show
which features were the most important when making a prediction'''

if abbrev in ['NB']:
    feature_order = np.argsort(np.mean(np.abs(shap_values.values), axis=0))
    df = pd.DataFrame({"Features": [feature_names[i] for i in feature_order][::-1], "Importance": [ np.mean(np.abs(

elif abbrev in ['LR', 'LGB', 'XGB', 'CGB']: #LR cant use shap_values.values
    feature_order = np.argsort(np.mean(np.abs(shap_values), axis=0))
    df = pd.DataFrame({"Features": [feature_names[i] for i in feature_order][::-1], "Importance": [ np.mean(np.abs(

else: # For multiclass models (can be used for NB)..Loops through Class 0 and Class 1
    # Sums up the shap average values form both classes to get the shap average for the whole CV for the model

    c_idx = []
    columns = feature_names
    for column in range(0, (len(columns))):
        if isinstance(shap_values, list):
            c_idx.append(X.columns.get_loc(column))
            means = [np.abs(shap_values[class_][:, c_idx]).mean(axis=0) for class_ in range(len(shap_values))]
            shap_means = np.sum(np.column_stack(means), 1)
        else:
            # Else there is only one 2D array of shap values
            assert len(shap_values.shape) == 2, 'Expected two-dimensional shap values array.'
            shap_means = np.abs(shap_values).mean(axis=0)
    df = pd.DataFrame({'Features': feature_names, 'Importance': shap_means}).sort_values(by='Importance', ascending=True)
    df.index += 1

return df
```

```
In [518... def merge_shap_df(abbrev, filepath, feat_order_map): # 'df' parameter is the dataframe returned from shap_feature_rank
'''Create a new dataframe that stores the model's shap feature importance values over each CV
and combines with features from original dataset'''

os.chdir(filepath)
extension = 'csv'

all_filenames = [i for i in glob.glob(abbrev+'_*.{}'.format(extension))]
print(all_filenames)

LR_shapFI = pd.concat([pd.read_csv(f) for f in all_filenames ])
print(LR_shapFI)
LR_shapFI.to_csv(abbrev + "_shapFI.csv", index=False, encoding='utf-8-sig')
# files = (filepath + 'LR_0_shapFIValues_Test.csv', filepath + 'LR_1_shapFIValues_Test.csv')

# result = pd.DataFrame()
# temp = result
# for f in files:

# df = pd.read_csv(f, sep=',', index_col=[0]).set_index('Features')
# temp = df[:-1]
# # print(type(temp))
# result = pd.merge(df, temp, on=['Features'], how='right')
# del(temp)
# # print(result)

# result.reset_index(inplace=True)

# return result
```

```
In [519... original_headers = pd.read_csv(full_path+"/exploratory/OriginalFeatureNames.csv",sep=',').columns.values.tolist() #Get
feat_order_map = {feat:i for i, feat in enumerate(original_headers)}

full_path = experiment_path+'/hcc-data_example'
filepath = full_path+"/model_evaluation/shap_values/testResults/" #path to save SHAP FI value results

# for cvCount in range(0, cv_partitions):
merge_shap_df('LR', filepath, feat_order_map)

['LR_0_shapFIValues_Test.csv', 'LR_2_shapFIValues_Test.csv', 'LR_shapFI.csv', 'LR_1_shapFIValues_Test.csv']
  Unnamed: 0      Features  Importance
0          0  Performance Status*    0.066918
1          1    Liver Metastasis    0.058906
2          2  Portal Vein Thrombosis    0.050533
3          3          Diabetes    0.044112
4          4    Esophageal Varices    0.043612
..      ...      ...      ...
36         36  Hepatitis B e Antigen    0.011481
37         37  Hepatitis B Surface Antigen    0.011007
38         38    Oxygen Saturation (%)    0.009104
39         39    Liver Metastasis    0.006661
40         40          Alcohol    0.004921

[232 rows x 3 columns]
```

Next Steps

- Saving shap figures per model in each cv
- Make sure you can loop through each pickled model, load it, create shap values and display plots
- Be able to load one model at a time, create shapley values for each CV train and test set, store shap scores in a dataframe
- Make sure to load original dataset features so that each csv file is the same length as the original dataset
 - This means when a CV dataset is missing a feature, we make sure to assign a shap score of 0
 - each new csv file for loading shap scores of each trained model must include all features

```
LR_shap_all_CVs.csv ==>
LR_0 --> CV0
LR_1 --> CV1
LR_2 --> CV2
```

- Save dataframe for each model in a csv file

More Updates/Fixes

8/02/22

- Currently unsure if creating DataFrame for each model's shap_values should be done in compute_shap_values() or within the nested for-loop in testing cell

8/04/22

- Can create DataFrames for each CV but feature names most likely are not matching actual values (double check it)
- Difficult looping through to merge Dataframes for all CVs features...tried temporary variable
- Must also consider that shap_values array are returned in order of features from test/train set it was passed from...not based on feature order in test/train set **FIXED on 8/05/22**
 - Consider mapping out and ordering the values to avoid shuffling of names and values **FIXED on 8/05/22**

8/05/22

- Saving feature importance scores for each cv
- Created two different runs, one for actual test (default) and another if the user chooses to run it on the training sets for comparison

8/08/22

- Iterating through multiclass shap values for Decision Tree poses issue?...ideally we'd want to get the shap absolute average for both classes 0 and 1...same might be for XGB and any other model that has multiclass output **FIXED on 8/08/22**
 - Figured out that when running the loop in shap_feature_ranking() for Decision Tree, both classes 0 and 1 are accounted for. The shap absolute averages are summed up automatically to get the overall CV feature importances for the model (i double checked this myself through creating a loop that would output two different csv files for each class it iterated through)
- **Current issue:** Figuring out how to save multiple figures for each model when calling shap_summary()...for now, I can only save each figure individually through each CV...if model NB has 2 plot function calls & iterate through 3 CVs --> total 6 shap plots for **ONE** model.....
 - **POSSIBLE FIX** merge all images onto one pdf per model which would entail different shap summaries **OR** create the master list of feature impmortance of all CVs for each model and create shap summaries for those

Run SHAP for Testing Datasets

Loop through each hcc_demo dataset to unpickle and load trained models to create Shapley values and plots Default run

- The default setting runs explainer and shap values for the TESTING datasets for each model and CV
- User has the option below to run the loop for training sets as well

```
In [18]: # testing all methods
run_force_plots = False # parameter in run_force_plot(); set to True if user wants to display force plots for trained
run_test = True
save_path = experiment_path + '/hcc-data_example/model_evaluation/shap_values/testResults/shapFigures/'

if run_test == True:
    for each in datasets:
        print("-----")
        print(each)
        print("-----")
        full_path = experiment_path+'/'+ each
        filepath = full_path+"/model_evaluation/shap_values/testResults/" #path to save SHAP FI value results

        #Make folder in experiment folder/datafolder to store all shap_values per algorithm/CV combination
        if not os.path.exists(full_path+'/model_evaluation/shap_values/testResults'):
            os.mkdir(full_path+'/model_evaluation/shap_values/testResults')

        original_headers = pd.read_csv(full_path+"/exploratory/OriginalFeatureNames.csv",sep=',').columns.values.tolist
```



```

feat_order_map = {feat:i for i, feat in enumerate(original_headers)}
print(feat_order_map)

for algorithm in algorithms: #loop through algorithms
    print(abbrev[algorithm])

    for cvCount in range(0,cv_partitions): #loop through cv's
        print('{}{} In CV{}...'.format(abbrev[algorithm], cvCount, cvCount))

        # unpickle and load model
        result_file = full_path+ '/models/pickledModels/' + abbrev[algorithm]+ "_" + str(cvCount)+".pickle"
        file = open(result_file, 'rb')
        model = pickle.load(file)
        file.close()
        print('\nChecking if correct model is loaded...\n', model)

        # Load CV datasets, paths to datasets updates with each iteration
        train_path = experiment_path + '/' + each + '/CVDatasets/' + '/' + each + '_CV_' + str(cvCount) + '_Tr
        test_path = experiment_path + '/' + each + '/CVDatasets/' + '/' + each + '_CV_' + str(cvCount) + '_Test.
        trainX, trainY, testX, testY, train_feat, test_feat = dataPrep(train_path,instance_label,class_label, t

        # shap computation and plots
        # Sanity check: print explainer to check if explainer exists
        explainer = get_explainer(model, abbrev[algorithm], trainX) #explainer must always use training set
        print('\nChecking explainer for {}{}...\n{}'.format(abbrev[algorithm], cvCount, explainer))

        print('\nChecking shap values for {}{}...\n'.format(abbrev[algorithm], cvCount))
        shap_values = compute_shapValues(model, abbrev[algorithm], explainer, testX)

        print('\nChecking shap plots for {}{}...\n'.format(abbrev[algorithm], cvCount))
        shap_summary(abbrev[algorithm], test_feat, shap_values, explainer, testX, cvCount)

        #save SHAP FI results for each model per CV
        print('\nChecking feature importance for {}{}...\n'.format(abbrev[algorithm], cvCount))
        shap_fi_df = shap_feature_ranking(abbrev[algorithm], shap_values, testX, test_feat) # can either choos
        shapFI_path = filepath+ abbrev[algorithm] + '_' + str(cvCount) + "_shapFIValues_Test.csv"
        shap_fi_df.to_csv(shapFI_path, header=True, index=True)

        #create new DataFrame of SHAP FI values from each CV with features from original dataset
        # merge_shap_df(abbrev[algorithm], filepath, feat_order_map, cvCount)
        # files = os.path.join(filepath, abbrev[algorithm] + '_' + str(cvCount) + "*.csv")
        # print('\nChecking for list of csv files...\n')
        # csv_files = glob.glob(files)
        # print(csv_files)

        # create new folder to save summary plots for each model per CV
        # if not os.path.exists(experiment_path+'/hcc-data_example/model_evaluation/shap_values/testResults/sh
        # os.mkdir(full_path+'/model_evaluation/shap_values/testResults/shapFigures')

        # filepath2 = full_path+"/model_evaluation/shap_values/testResults/shapFigures"+ abbrev[algorithm]
        # summary.to_pdf(filepath2, header=True, index=True)

        # only runs force plots if run = True
        # if run_force_plots == True:
        #     if abbrev[algorithm] in ['NB']:

        #         print('\nForce Plot for {}{} SHAP Values in Test Set: \n'.format(abbrev[algorithm], cvCount))
        #         shap.force_plot(shap_values, feature_names = test_feat)

        #         print('\nSingle-Prediction Force Plot for {}{} SHAP Values in Test Set: \n'.format(abbrev[al
        #         shap.force_plot(shap_values[42], testX.iloc[42], feature_names=test_feat, show=True)
        #         break

        #     elif abbrev[algorithm] in ['LR', 'XGB', 'LGB', 'CBG']: #need to test out LGB and CBG for this

        #         print('\nForce Plot for {}{} SHAP Values in Whole Test Set: \n'.format(abbrev[algorithm], c
        #         shap.force_plot(explainer.expected_value, shap_values, testX, feature_names=test_feat)

        #         print('\nSingle-Prediction Force Plot for {}{} SHAP Values in Test Set: \n'.format(abbrev[al
        #         shap.force_plot(explainer.expected_value, shap_values[42], testX.iloc[42], feature_names=tes
        #         break

        #     else:
        #         # Decision Tree has multiclass output so needed to create two separate function calls
        #         # Decision Tree doesn't work when just using shap_values as a parameter
        #         print('\nForce Plot for {}{} SHAP Values from Class 0 in Test Set: \n'.format(abbrev[algorit
        #         shap.force_plot(explainer.expected_value[0], shap_values[0], feature_names=test_feat)

        #         print('\nForce Plot for {}{} SHAP Values from Class 1 in Test Set: \n'.format(abbrev[algorit
        #         shap.force_plot(explainer.expected_value[1], shap_values[1], feature_names=test_feat)
        #         break

```

hcc-data_example

```
{'Gender': 0, 'Symptoms ': 1, 'Alcohol': 2, 'Hepatitis B Surface Antigen': 3, 'Hepatitis B e Antigen': 4, 'Hepatitis B Core Antibody': 5, 'Hepatitis C Virus Antibody': 6, 'Cirrhosis': 7, 'Endemic Countries': 8, 'Smoking': 9, 'Diabetes': 10, 'Obesity': 11, 'Hemochromatosis': 12, 'Arterial Hypertension': 13, 'Chronic Renal Insufficiency': 14, 'Human Immunodeficiency Virus': 15, 'Nonalcoholic Steatohepatitis': 16, 'Esophageal Varices': 17, 'Splenomegaly': 18, 'Portal Hypertension': 19, 'Portal Vein Thrombosis': 20, 'Liver Metastasis': 21, 'Radiological Hallmark': 22, 'Age at diagnosis': 23, 'Grams of Alcohol per day': 24, 'Packs of cigarets per year': 25, 'Performance Status*': 26, 'Encephalopathy degree*': 27, 'Ascites degree*': 28, 'International Normalised Ratio*': 29, 'Alpha-Fetoprotein (ng/mL)': 30, 'Haemoglobin (g/dL)': 31, 'Mean Corpuscular Volume': 32, 'Leukocytes(G/L)': 33, 'Platelets': 34, 'Albumin (mg/dL)': 35, 'Total Bilirubin(mg/dL)': 36, 'Alanine transaminase (U/L)': 37, 'Aspartate transaminase (U/L)': 38, 'Gamma glutamyl transferase (U/L)': 39, 'Alkaline phosphatase (U/L)': 40, 'Total Proteins (g/dL)': 41, 'Creatinine (mg/dL)': 42, 'Number of Nodules': 43, 'Major dimension of nodule (cm)': 44, 'Direct Bilirubin (mg/dL)': 45, 'Iron': 46, 'Oxygen Saturation (%)': 47, 'Ferritin (ng/mL)': 48}
NB
NB0 In CV0...
```

Checking if correct model is loaded...

```
GaussianNB()
['Alanine transaminase (U/L)', 'Albumin (mg/dL)', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/mL)', 'Arterial Hypertension', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Chronic Renal Insufficiency', 'Cirrhosis', 'Creatinine (mg/dL)', 'Diabetes', 'Direct Bilirubin (mg/dL)', 'Encephalopathy degree*', 'Endemic Countries', 'Esophageal Varices', 'Ferritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haemoglobin (g/dL)', 'Hemochromatosis', 'Hepatitis B Surface Antigen', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibody', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Number of Nodules', 'Obesity', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Portal Hypertension', 'Portal Vein Thrombosis', 'Smoking', 'Splenomegaly', 'Symptoms ', 'Total Bilirubin(mg/dL)']
['Alanine transaminase (U/L)', 'Albumin (mg/dL)', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/mL)', 'Arterial Hypertension', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Chronic Renal Insufficiency', 'Cirrhosis', 'Creatinine (mg/dL)', 'Diabetes', 'Direct Bilirubin (mg/dL)', 'Encephalopathy degree*', 'Endemic Countries', 'Esophageal Varices', 'Ferritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haemoglobin (g/dL)', 'Hemochromatosis', 'Hepatitis B Surface Antigen', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibody', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Number of Nodules', 'Obesity', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Portal Hypertension', 'Portal Vein Thrombosis', 'Smoking', 'Splenomegaly', 'Symptoms ', 'Total Bilirubin(mg/dL)']
```

Checking explainer for NB0...

```
shap.explainers.Permutation()
```

Checking shap values for NB0...

```
.values =
array([[ 5.00000000e-03, -3.16666667e-02, -1.91666667e-02, ...,
        -2.50000000e-03,  4.16666667e-03, -1.75000000e-02],
       [-1.66666667e-03,  2.50000000e-02,  2.66666667e-02, ...,
         9.16666667e-03,  3.75000000e-02, -2.83333333e-02],
       [ 2.50000000e-03,  2.31296463e-18,  1.50000000e-02, ...,
         2.50000000e-03,  3.16666667e-02, -9.16666667e-03],
       ...,
       [ 1.66666667e-03, -1.16666667e-02, -8.33333333e-03, ...,
         1.66666667e-03,  3.33333333e-03, -2.58333333e-02],
       [-4.16666667e-03, -2.50000000e-03, -1.33333333e-02, ...,
         0.00000000e+00, -3.00000000e-02, -2.83333333e-02],
       [ 8.33333333e-03,  3.16666667e-02, -1.25000000e-02, ...,
         8.33333333e-04, -1.15000000e-01,  4.90000000e-01]])

.base_values =
array([0.33, 0.33, 0.33, 0.33, 0.33, 0.33, 0.33, 0.33, 0.33, 0.33, 0.33,
       0.33, 0.33, 0.33, 0.33, 0.33, 0.33, 0.33, 0.33, 0.33, 0.33, 0.33,
       0.33, 0.33, 0.33, 0.33, 0.33, 0.33, 0.33, 0.33, 0.33, 0.33, 0.33,
       0.33, 0.33, 0.33, 0.33, 0.33, 0.33, 0.33, 0.33, 0.33, 0.33, 0.33])

.data =
array([[ 0.0368995,  1.0551286, -0.5852099, ..., -1.2247449,  0.6264224,
        -0.4383745],
       [-0.3323658, -0.2719725,  3.8874108, ...,  0.8164966,  0.6264224,
        -0.3900995],
       [-0.0973788,  0.3178502,  0.1974987, ...,  0.8164966,  0.6264224,
        -0.4866495],
       ...,
       [-0.6680615,  0.6127615, -0.708207 , ...,  0.8164966,  0.6264224,
        -0.3740079],
       [-0.9198333,  0.3178502, -0.484576 , ...,  0.8164966, -1.5963668,
        -0.2613663],
       [ 0.1376082, -0.3457004, -0.3615789, ...,  0.8164966, -1.5963668,
         1.1546993]])
```

Checking shap plots for NB0...

Summary Plot for SHAP Values in Class 0 & 1 in Test Set:

SHAP Beeswarm Plot for Top 5 SHAP Values in Class 0 & 1 in Test Set:

Checking feature importance for NB0...

Checking for list of csv files...

```
['/Users/jessicakim/Desktop/STREAMLINE/DemoData/Output/hcc_demo/hcc-data_example/model_evaluation/shap_values/testResults/NB_0_shapFIValues_Test.csv']
```

Unnamed: 0		Features	Importance
0	0	Creatinine (mg/dL)	0.073712
1	1	Encephalopathy degree*	0.059318
2	2	Total Bilirubin(mg/dL)	0.044727
3	3	Performance Status*	0.037212
4	4	Direct Bilirubin (mg/dL)	0.036152
5	5	Liver Metastasis	0.035000
6	6	Hemochromatosis	0.034303
7	7	Aspartate transaminase (U/L)	0.032030
8	8	Ascites degree*	0.029288
9	9	Chronic Renal Insufficiency	0.029273
10	10	Portal Vein Thrombosis	0.027273
11	11	Packs of cigarets per year	0.026742
12	12	Iron	0.024727
13	13	Obesity	0.024424
14	14	International Normalised Ratio*	0.023924
15	15	Major dimension of nodule (cm)	0.023682
16	16	Symptoms	0.023106
17	17	Esophageal Varices	0.022485
18	18	Alpha-Fetoprotein (ng/mL)	0.020182
19	19	Albumin (mg/dL)	0.018879
20	20	Endemic Countries	0.017348
21	21	Oxygen Saturation (%)	0.015091
22	22	Diabetes	0.014273
23	23	Haemoglobin (g/dL)	0.014227
24	24	Hepatitis C Virus Antibody	0.013879
25	25	Alkaline phosphatase (U/L)	0.013636
26	26	Hepatitis B Surface Antigen	0.012864
27	27	Leukocytes(G/L)	0.012773
28	28	Gamma glutamyl transferase (U/L)	0.011833
29	29	Ferritin (ng/mL)	0.009318
30	30	Cirrhosis	0.008485
31	31	Mean Corpuscular Volume	0.005955
32	32	Alanine transaminase (U/L)	0.004530
33	33	Number of Nodules	0.003364
34	34	Smoking	0.002515
35	35	Arterial Hypertension	0.002394
36	36	Portal Hypertension	0.002015
37	37	Splenomegaly	0.001939
38	38	Hepatitis B e Antigen	0.000000

Out [18]:

Unnamed: 0		Features	Importance
0	0	Creatinine (mg/dL)	0.073712
1	1	Encephalopathy degree*	0.059318
2	2	Total Bilirubin(mg/dL)	0.044727
3	3	Performance Status*	0.037212
4	4	Direct Bilirubin (mg/dL)	0.036152
5	5	Liver Metastasis	0.035000
6	6	Hemochromatosis	0.034303
7	7	Aspartate transaminase (U/L)	0.032030
8	8	Ascites degree*	0.029288
9	9	Chronic Renal Insufficiency	0.029273
10	10	Portal Vein Thrombosis	0.027273
11	11	Packs of cigarets per year	0.026742
12	12	Iron	0.024727
13	13	Obesity	0.024424
14	14	International Normalised Ratio*	0.023924
15	15	Major dimension of nodule (cm)	0.023682
16	16	Symptoms	0.023106
17	17	Esophageal Varices	0.022485
18	18	Alpha-Fetoprotein (ng/mL)	0.020182
19	19	Albumin (mg/dL)	0.018879
20	20	Endemic Countries	0.017348
21	21	Oxygen Saturation (%)	0.015091
22	22	Diabetes	0.014273
23	23	Haemoglobin (g/dL)	0.014227
24	24	Hepatitis C Virus Antibody	0.013879
25	25	Alkaline phosphatase (U/L)	0.013636
26	26	Hepatitis B Surface Antigen	0.012864
27	27	Leukocytes(G/L)	0.012773
28	28	Gamma glutamyl transferase (U/L)	0.011833
29	29	Ferritin (ng/mL)	0.009318
30	30	Cirrhosis	0.008485
31	31	Mean Corpuscular Volume	0.005955
32	32	Alanine transaminase (U/L)	0.004530
33	33	Number of Nodules	0.003364
34	34	Smoking	0.002515
35	35	Arterial Hypertension	0.002394
36	36	Portal Hypertension	0.002015
37	37	Splenomegaly	0.001939
38	38	Hepatitis B e Antigen	0.000000

NB1 In CV1...

```
Checking if correct model is loaded...
GaussianNB()
['Age at diagnosis', 'Alanine transaminase (U/L)', 'Albumin (mg/dL)', 'Alcohol', 'Alkaline phosphatase (U/L)', 'Alpha-
Fetoprotein (ng/mL)', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Cirrhosis', 'Creatinine (mg/dL)', 'Diabete
s', 'Direct Bilirubin (mg/dL)', 'Endemic Countries', 'Esophageal Varices', 'Ferritin (ng/mL)', 'Gamma glutamyl transfe
rase (U/L)', 'Grams of Alcohol per day', 'Haemoglobin (g/dL)', 'Hepatitis B Core Antibody', 'Hepatitis B Surface Antig
en', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibody', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/
L)', 'Liver Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Nonalcoholic Steatohepatitis',
'Number of Nodules', 'Obesity', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Platele
ts', 'Portal Hypertension', 'Portal Vein Thrombosis', 'Smoking', 'Symptoms ', 'Total Bilirubin(mg/dL)', 'Total Protein
s (g/dL)']
['Age at diagnosis', 'Alanine transaminase (U/L)', 'Albumin (mg/dL)', 'Alcohol', 'Alkaline phosphatase (U/L)', 'Alpha-
Fetoprotein (ng/mL)', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Cirrhosis', 'Creatinine (mg/dL)', 'Diabete
s', 'Direct Bilirubin (mg/dL)', 'Endemic Countries', 'Esophageal Varices', 'Ferritin (ng/mL)', 'Gamma glutamyl transfe
rase (U/L)', 'Grams of Alcohol per day', 'Haemoglobin (g/dL)', 'Hepatitis B Core Antibody', 'Hepatitis B Surface Antig
en', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibody', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/
L)', 'Liver Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Nonalcoholic Steatohepatitis',
'Number of Nodules', 'Obesity', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Platele
ts', 'Portal Hypertension', 'Portal Vein Thrombosis', 'Smoking', 'Symptoms ', 'Total Bilirubin(mg/dL)', 'Total Protein
s (g/dL)']

Checking explainer for NB1...
shap.explainers.Permutation()

Checking shap values for NB1...

.values =
array([[ -0.01416667,  0.025      , -0.10666667, ...,  0.12166667,
        -0.0175      , -0.01166667],
       [ 0.03333333, -0.00833333, -0.01166667, ..., -0.2925      ,
        -0.01833333, -0.02        ],
       [ 0.01583333,  0.01166667,  0.01666667, ..., -0.04166667,
        -0.01333333, -0.00916667],
       ...,
       [-0.01666667, -0.01333333,  0.02416667, ..., -0.09166667,
        -0.02333333, -0.01416667],
       [-0.00083333, -0.00833333, -0.00083333, ...,  0.01916667,
        -0.00166667, -0.00333333],
       [ 0.03666667,  0.0175      , -0.04416667, ...,  0.10333333,
        -0.01666667, -0.01        ]])

.base_values =
array([0.49, 0.49, 0.49, 0.49, 0.49, 0.49, 0.49, 0.49, 0.49, 0.49, 0.49,
       0.49, 0.49, 0.49, 0.49, 0.49, 0.49, 0.49, 0.49, 0.49, 0.49, 0.49,
       0.49, 0.49, 0.49, 0.49, 0.49, 0.49, 0.49, 0.49, 0.49, 0.49, 0.49,
       0.49, 0.49, 0.49, 0.49, 0.49, 0.49, 0.49, 0.49, 0.49, 0.49, 0.49,
       0.49, 0.49, 0.49, 0.49, 0.49, 0.49, 0.49, 0.49, 0.49, 0.49])

.data =
array([[ -0.7178553, -0.3317865,  1.9018142, ...,  0.6831301, -0.4188214,
        -0.0949777],
       [ 0.9544964, -0.9389338,  0.0448948, ..., -1.4638501, -0.4505005,
        -0.2135745],
       [ 0.5909417, -0.2169208, -0.558604 , ..., -1.4638501, -0.1812288,
        -0.176513 ],
       ...,
       [-0.4270115,  0.83328   ,  0.1996381, ..., -1.4638501, -0.3713029,
        -0.2358114],
       [-0.4270115,  1.1450584, -0.1098485, ...,  0.6831301, -0.2445869,
        -0.2358114],
       [ 1.0272073, -0.6763836,  1.1280978, ...,  0.6831301, -0.3871424,
        -0.2580483]])

Checking shap plots for NB1...

Summary Plot for SHAP Values in Class 0 & 1 in Test Set:

SHAP Beeswarm Plot for Top 5 SHAP Values in Class 0 & 1 in Test Set:
```

Checking feature importance for NB1...

```
Checking for list of csv files...

['/Users/jessicakim/Desktop/STREAMLINE/DemoData/Output/hcc_demo/hcc-data_example/model_evaluation/shap_values/testResu
lts/NB_1_shapFIValues_Test.csv']
  Unnamed: 0      Features  Importance
0           0      Endemic Countries  0.099591
1           1  Nonalcoholic Steatohepatitis  0.068121
2           2      Symptoms  0.063303
3           3  Performance Status*  0.058742
4           4      Ascites degree*  0.055061
5           5  Alkaline phosphatase (U/L)  0.047879
6           6      Albumin (mg/dL)  0.043242
7           7  Portal Hypertension  0.040152
8           8  International Normalised Ratio*  0.039545
9           9      Haemoglobin (g/dL)  0.038909
10          10      Total Bilirubin(mg/dL)  0.029288
11          11  Alpha-Fetoprotein (ng/mL)  0.028136
12          12  Portal Vein Thrombosis  0.028061
```


13	13	Liver Metastasis	0.027333
14	14	Grams of Alcohol per day	0.025697
15	15	Oxygen Saturation (%)	0.023758
16	16	Iron	0.023439
17	17	Direct Bilirubin (mg/dL)	0.021667
18	18	Platelets	0.018500
19	19	Age at diagnosis	0.018439
20	20	Alanine transaminase (U/L)	0.015803
21	21	Mean Corpuscular Volume	0.015697
22	22	Packs of cigarets per year	0.015242
23	23	Major dimension of nodule (cm)	0.015182
24	24	Ferritin (ng/mL)	0.014803
25	25	Hepatitis B Surface Antigen	0.014136
26	26	Total Proteins (g/dL)	0.013788
27	27	Obesity	0.011015
28	28	Diabetes	0.010985
29	29	Aspartate transaminase (U/L)	0.008742
30	30	Hepatitis C Virus Antibody	0.008515
31	31	Alcohol	0.006924
32	32	Hepatitis B Core Antibody	0.006742
33	33	Gamma glutamyl transferase (U/L)	0.006152
34	34	Smoking	0.005591
35	35	Number of Nodules	0.003652
36	36	Creatinine (mg/dL)	0.003515
37	37	Esophageal Varices	0.003061
38	38	Hepatitis B e Antigen	0.002636
39	39	Leukocytes(G/L)	0.000258
40	40	Cirrhosis	0.000000

Out [18] :

Unnamed: 0		Features	Importance
0	0	Endemic Countries	0.099591
1	1	Nonalcoholic Steatohepatitis	0.068121
2	2	Symptoms	0.063303
3	3	Performance Status*	0.058742
4	4	Ascites degree*	0.055061
5	5	Alkaline phosphatase (U/L)	0.047879
6	6	Albumin (mg/dL)	0.043242
7	7	Portal Hypertension	0.040152
8	8	International Normalised Ratio*	0.039545
9	9	Haemoglobin (g/dL)	0.038909
10	10	Total Bilirubin(mg/dL)	0.029288
11	11	Alpha-Fetoprotein (ng/mL)	0.028136
12	12	Portal Vein Thrombosis	0.028061
13	13	Liver Metastasis	0.027333
14	14	Grams of Alcohol per day	0.025697
15	15	Oxygen Saturation (%)	0.023758
16	16	Iron	0.023439
17	17	Direct Bilirubin (mg/dL)	0.021667
18	18	Platelets	0.018500
19	19	Age at diagnosis	0.018439
20	20	Alanine transaminase (U/L)	0.015803
21	21	Mean Corpuscular Volume	0.015697
22	22	Packs of cigarets per year	0.015242
23	23	Major dimension of nodule (cm)	0.015182
24	24	Ferritin (ng/mL)	0.014803
25	25	Hepatitis B Surface Antigen	0.014136
26	26	Total Proteins (g/dL)	0.013788
27	27	Obesity	0.011015
28	28	Diabetes	0.010985
29	29	Aspartate transaminase (U/L)	0.008742
30	30	Hepatitis C Virus Antibody	0.008515
31	31	Alcohol	0.006924
32	32	Hepatitis B Core Antibody	0.006742
33	33	Gamma glutamyl transferase (U/L)	0.006152
34	34	Smoking	0.005591
35	35	Number of Nodules	0.003652
36	36	Creatinine (mg/dL)	0.003515
37	37	Esophageal Varices	0.003061
38	38	Hepatitis B e Antigen	0.002636
39	39	Leukocytes(G/L)	0.000258
40	40	Cirrhosis	0.000000

NB2 In CV2...

```
Checking if correct model is loaded...
GaussianNB()
['Age at diagnosis', 'Albumin (mg/dL)', 'Alcohol', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/mL)', 'Arterial Hypertension', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Chronic Renal Insufficiency', 'Creatinine (mg/dL)', 'Diabetes', 'Direct Bilirubin (mg/dL)', 'Encephalopathy degree*', 'Endemic Countries', 'Ferritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haemoglobin (g/dL)', 'Hepatitis B Core Antibody', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibody', 'Human Immunodeficiency Virus', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Number of Nodules', 'Obesity', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Platelets', 'Portal Hypertension', 'Portal Vein Thrombosis', 'Symptoms ']
```

Checking explainer for NB2...
shap.explainers.Permutation()

Checking shap values for NB2...

```
.values =
array([[ 0.          ,  0.          ,  0.          , ...,  0.          ,
         0.          , -0.0025       ],
       [ 0.00416667,  0.00083333,  0.          , ...,  0.          ,
         0.          ,  0.          ],
       [ 0.01333333,  0.00333333, -0.00333333, ..., -0.00583333,
        -0.00583333,  0.0025       ],
       ...,
       [ 0.          ,  0.01833333,  0.          , ..., -0.0025       ,
        -0.00416667,  0.005        ],
       [-0.0025      , -0.05916667,  0.00916667, ..., -0.00583333,
        -0.03916667,  0.00916667],
       [ 0.          ,  0.          ,  0.          , ...,  0.          ,
         0.          ,  0.          ]])

.base_values =
array([0.05, 0.05, 0.05, 0.05, 0.05, 0.05, 0.05, 0.05, 0.05, 0.05, 0.05,
       0.05, 0.05, 0.05, 0.05, 0.05, 0.05, 0.05, 0.05, 0.05, 0.05, 0.05,
       0.05, 0.05, 0.05, 0.05, 0.05, 0.05, 0.05, 0.05, 0.05, 0.05, 0.05,
       0.05, 0.05, 0.05, 0.05, 0.05, 0.05, 0.05, 0.05, 0.05, 0.05])

.data =
array([[ -0.8437697, -0.4184704,  0.5703518, ...,  0.6264224, -0.5282705,
        -1.3228757],
       [ -3.7262129, -0.4184704,  0.5703518, ..., -1.5963668, -0.5282705,
         0.7559289],
       [ -1.0908363,  0.0205566, -1.7533038, ...,  0.6264224, -0.5282705,
         0.7559289],
       ...,
       [ -1.0908363, -1.8818937,  0.5703518, ...,  0.6264224, -0.5282705,
         0.7559289],
       [ -1.2555473,  1.0449529,  0.5703518, ...,  0.6264224, -0.5282705,
         0.7559289],
       [ -0.1025701, -2.028236 ,  0.5703518, ...,  0.6264224, -0.5282705,
         0.7559289]])
```

Checking shap plots for NB2...

Summary Plot for SHAP Values in Class 0 & 1 in Test Set:

SHAP Beeswarm Plot for Top 5 SHAP Values in Class 0 & 1 in Test Set:

Checking feature importance for NB2...

Checking for list of csv files...

```
['/Users/jessicakim/Desktop/STREAMLINE/DemoData/Output/hcc_demo/hcc-data_example/model_evaluation/shap_values/testResults/NB_2_shapFIValues_Test.csv']
```

Unnamed: 0	Features	Importance
0	Direct Bilirubin (mg/dL)	0.054727
1	Human Immunodeficiency Virus	0.053182
2	Packs of cigarets per year	0.017985
3	Ferritin (ng/mL)	0.015485
4	International Normalised Ratio*	0.014712
5	Hepatitis B e Antigen	0.009273
6	Liver Metastasis	0.006045
7	Leukocytes(G/L)	0.004591
8	Portal Vein Thrombosis	0.003530
9	Obesity	0.003333
10	Ascites degree*	0.003015
11	Performance Status*	0.003000
12	Aspartate transaminase (U/L)	0.002985
13	Haemoglobin (g/dL)	0.002788
14	Alkaline phosphatase (U/L)	0.002758

15	15	Hepatitis B Core Antibody	0.002485
16	16	Alpha-Fetoprotein (ng/mL)	0.002000
17	17	Encephalopathy degree*	0.001909
18	18	Albumin (mg/dL)	0.001909
19	19	Creatinine (mg/dL)	0.001758
20	20	Platelets	0.001515
21	21	Number of Nodules	0.001470
22	22	Diabetes	0.001439
23	23	Major dimension of nodule (cm)	0.001424
24	24	Mean Corpuscular Volume	0.001242
25	25	Hepatitis C Virus Antibody	0.000879
26	26	Age at diagnosis	0.000879
27	27	Gamma glutamyl transferase (U/L)	0.000848
28	28	Iron	0.000667
29	29	Chronic Renal Insufficiency	0.000561
30	30	Symptoms	0.000530
31	31	Portal Hypertension	0.000515
32	32	Arterial Hypertension	0.000500
33	33	Alcohol	0.000455
34	34	Oxygen Saturation (%)	0.000212
35	35	Endemic Countries	0.000091

Out [18]:

Unnamed: 0		Features	Importance
0	0	Direct Bilirubin (mg/dL)	0.054727
1	1	Human Immunodeficiency Virus	0.053182
2	2	Packs of cigarets per year	0.017985
3	3	Ferritin (ng/mL)	0.015485
4	4	International Normalised Ratio*	0.014712
5	5	Hepatitis B e Antigen	0.009273
6	6	Liver Metastasis	0.006045
7	7	Leukocytes(G/L)	0.004591
8	8	Portal Vein Thrombosis	0.003530
9	9	Obesity	0.003333
10	10	Ascites degree*	0.003015
11	11	Performance Status*	0.003000
12	12	Aspartate transaminase (U/L)	0.002985
13	13	Haemoglobin (g/dL)	0.002788
14	14	Alkaline phosphatase (U/L)	0.002758
15	15	Hepatitis B Core Antibody	0.002485
16	16	Alpha-Fetoprotein (ng/mL)	0.002000
17	17	Encephalopathy degree*	0.001909
18	18	Albumin (mg/dL)	0.001909
19	19	Creatinine (mg/dL)	0.001758
20	20	Platelets	0.001515
21	21	Number of Nodules	0.001470
22	22	Diabetes	0.001439
23	23	Major dimension of nodule (cm)	0.001424
24	24	Mean Corpuscular Volume	0.001242
25	25	Hepatitis C Virus Antibody	0.000879
26	26	Age at diagnosis	0.000879
27	27	Gamma glutamyl transferase (U/L)	0.000848
28	28	Iron	0.000667
29	29	Chronic Renal Insufficiency	0.000561
30	30	Symptoms	0.000530
31	31	Portal Hypertension	0.000515
32	32	Arterial Hypertension	0.000500
33	33	Alcohol	0.000455
34	34	Oxygen Saturation (%)	0.000212
35	35	Endemic Countries	0.000091

```
LR
LR0 In CV0...

Checking if correct model is loaded...
LogisticRegression(C=0.006606805070193189, dual=True,
                    max_iter=193.8544995971634, random_state=42,
                    solver='liblinear')
['Alanine transaminase (U/L)', 'Albumin (mg/dL)', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/mL)', 'Arterial Hypertension', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Chronic Renal Insufficiency', 'Cirrhosis', 'Creatinine (mg/dL)', 'Diabetes', 'Direct Bilirubin (mg/dL)', 'Encephalopathy degree*', 'Endemic Countries', 'Esophageal Varices', 'Ferritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haemoglobin (g/dL)', 'Hemochromatosis', 'Hepatitis B Surface Antigen', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibody', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Number of Nodules', 'Obesity', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Portal Hypertension', 'Portal Vein Thrombosis', 'Smoking', 'Splenomegaly', 'Symptoms ', 'Total Bilirubin(mg/dL)']
['Alanine transaminase (U/L)', 'Albumin (mg/dL)', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/mL)', 'Arterial Hypertension', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Chronic Renal Insufficiency', 'Cirrhosis', 'Creatinine (mg/dL)', 'Diabetes', 'Direct Bilirubin (mg/dL)', 'Encephalopathy degree*', 'Endemic Countries', 'Esophageal Varices', 'Ferritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haemoglobin (g/dL)', 'Hemochromatosis', 'Hepatitis B Surface Antigen', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibody', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Number of Nodules', 'Obesity', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Portal Hypertension', 'Portal Vein Thrombosis', 'Smoking', 'Splenomegaly', 'Symptoms ', 'Total Bilirubin(mg/dL)']
```

Checking explainer for LR0...
<shap.explainers._linear.Linear object at 0x7f88a6c09820>

Checking shap values for LR0...

```
[[ -1.54247265e-04 -5.20008470e-02 -4.40485958e-02 ... -1.47969848e-03
   2.31157863e-02 -2.02239904e-02]
 [  6.85778818e-04  1.94894821e-02  2.13041189e-01 ...  1.21066239e-03
   2.31157863e-02 -1.80842415e-02]
 [  1.51216786e-04 -1.22839969e-02  9.42115197e-04 ...  1.21066239e-03
   2.31157863e-02 -2.23637392e-02]
 ...
 [  1.44943883e-03 -2.81707337e-02 -5.11185666e-02 ...  1.21066239e-03
   2.31157863e-02 -1.73709948e-02]
 [  2.02218390e-03 -1.22839969e-02 -3.82640794e-02 ...  1.21066239e-03
  -6.24982370e-02 -1.23782505e-02]
 [-3.83345247e-04  2.34611703e-02 -3.11941086e-02 ...  1.21066239e-03
  -6.24982370e-02  5.03876684e-02]]
```

Checking shap plots for LR0...

Expected value for LR: -0.023696555525940875
Summary Plot for SHAP Values in Test Set:

SHAP Bar Plot for SHAP Values Test Set:

SHAP Decision Plot for SHAP Values in Test Set:

SHAP Decision Plot for Single-Prediction in Test Set:

Checking feature importance for LR0...

Checking for list of csv files...

```
['/Users/jessicakim/Desktop/STREAMLINE/DemoData/Output/hcc_demo/hcc-data_example/model_evaluation/shap_values/testResults/LR_0_shapFIValues_Test.csv']
```

Unnamed: 0	Features	Importance
0	Performance Status*	0.066918
1	Liver Metastasis	0.058906
2	Portal Vein Thrombosis	0.050533
3	Diabetes	0.044112
4	Esophageal Varices	0.043612
5	Ascites degree*	0.043348
6	Aspartate transaminase (U/L)	0.042648
7	Major dimension of nodule (cm)	0.042610
8	Albumin (mg/dL)	0.042516
9	Symptoms	0.038869
10	Alkaline phosphatase (U/L)	0.038658
11	Creatinine (mg/dL)	0.036096
12	Ferritin (ng/mL)	0.035923
13	Haemoglobin (g/dL)	0.033432
14	Gamma glutamyl transferase (U/L)	0.027842
15	Hepatitis C Virus Antibody	0.026838
16	Chronic Renal Insufficiency	0.021532
17	Portal Hypertension	0.021086
18	Leukocytes(G/L)	0.020526
19	Hepatitis B Surface Antigen	0.019027
20	Obesity	0.018688
21	Direct Bilirubin (mg/dL)	0.017843
22	Encephalopathy degree*	0.017500
23	International Normalised Ratio*	0.017237
24	Total Bilirubin(mg/dL)	0.016980
25	Endemic Countries	0.015877
26	Packs of cigarets per year	0.013487
27	Arterial Hypertension	0.010464
28	Cirrhosis	0.009670
29	Hemochromatosis	0.007440

30	30	Alpha-Fetoprotein (ng/mL)	0.006092
31	31	Iron	0.005802
32	32	Mean Corpuscular Volume	0.003474
33	33	Number of Nodules	0.002635
34	34	Oxygen Saturation (%)	0.002473
35	35	Smoking	0.002158
36	36	Alanine transaminase (U/L)	0.001406
37	37	Splenomegaly	0.001318
38	38	Hepatitis B e Antigen	0.000000

Out[18]:

Unnamed: 0		Features	Importance
0	0	Performance Status*	0.066918
1	1	Liver Metastasis	0.058906
2	2	Portal Vein Thrombosis	0.050533
3	3	Diabetes	0.044112
4	4	Esophageal Varices	0.043612
5	5	Ascites degree*	0.043348
6	6	Aspartate transaminase (U/L)	0.042648
7	7	Major dimension of nodule (cm)	0.042610
8	8	Albumin (mg/dL)	0.042516
9	9	Symptoms	0.038869
10	10	Alkaline phosphatase (U/L)	0.038658
11	11	Creatinine (mg/dL)	0.036096
12	12	Ferritin (ng/mL)	0.035923
13	13	Haemoglobin (g/dL)	0.033432
14	14	Gamma glutamyl transferase (U/L)	0.027842
15	15	Hepatitis C Virus Antibody	0.026838
16	16	Chronic Renal Insufficiency	0.021532
17	17	Portal Hypertension	0.021086
18	18	Leukocytes(G/L)	0.020526
19	19	Hepatitis B Surface Antigen	0.019027
20	20	Obesity	0.018688
21	21	Direct Bilirubin (mg/dL)	0.017843
22	22	Encephalopathy degree*	0.017500
23	23	International Normalised Ratio*	0.017237
24	24	Total Bilirubin(mg/dL)	0.016980
25	25	Endemic Countries	0.015877
26	26	Packs of cigarets per year	0.013487
27	27	Arterial Hypertension	0.010464
28	28	Cirrhosis	0.009670
29	29	Hemochromatosis	0.007440
30	30	Alpha-Fetoprotein (ng/mL)	0.006092
31	31	Iron	0.005802
32	32	Mean Corpuscular Volume	0.003474
33	33	Number of Nodules	0.002635
34	34	Oxygen Saturation (%)	0.002473
35	35	Smoking	0.002158
36	36	Alanine transaminase (U/L)	0.001406
37	37	Splenomegaly	0.001318
38	38	Hepatitis B e Antigen	0.000000

LR1 In CV1...

```
Checking if correct model is loaded...
LogisticRegression(C=0.06359900885943309, max_iter=48.076782938152924,
                    random_state=42, solver='sag')
['Age at diagnosis', 'Alanine transaminase (U/L)', 'Albumin (mg/dL)', 'Alcohol', 'Alkaline phosphatase (U/L)', 'Alpha-
Fetoprotein (ng/mL)', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Cirrhosis', 'Creatinine (mg/dL)', 'Diabete
s', 'Direct Bilirubin (mg/dL)', 'Endemic Countries', 'Esophageal Varices', 'Ferritin (ng/mL)', 'Gamma glutamyl transfe
rase (U/L)', 'Grams of Alcohol per day', 'Haemoglobin (g/dL)', 'Hepatitis B Core Antibody', 'Hepatitis B Surface Antig
en', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibody', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/
L)', 'Liver Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Nonalcoholic Steatohepatitis',
'Number of Nodules', 'Obesity', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Platele
ts', 'Portal Hypertension', 'Portal Vein Thrombosis', 'Smoking', 'Symptoms ', 'Total Bilirubin(mg/dL)', 'Total Protein
s (g/dL)']
['Age at diagnosis', 'Alanine transaminase (U/L)', 'Albumin (mg/dL)', 'Alcohol', 'Alkaline phosphatase (U/L)', 'Alpha-
Fetoprotein (ng/mL)', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Cirrhosis', 'Creatinine (mg/dL)', 'Diabete
s', 'Direct Bilirubin (mg/dL)', 'Endemic Countries', 'Esophageal Varices', 'Ferritin (ng/mL)', 'Gamma glutamyl transfe
rase (U/L)', 'Grams of Alcohol per day', 'Haemoglobin (g/dL)', 'Hepatitis B Core Antibody', 'Hepatitis B Surface Antig
en', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibody', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/
L)', 'Liver Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Nonalcoholic Steatohepatitis',
'Number of Nodules', 'Obesity', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Platele
ts', 'Portal Hypertension', 'Portal Vein Thrombosis', 'Smoking', 'Symptoms ', 'Total Bilirubin(mg/dL)', 'Total Protein
s (g/dL)']
```

Checking explainer for LR1...
<shap.explainers._linear.Linear object at 0x7f88c8b9d100>

Checking shap values for LR1...

```
[[[-0.19905492  0.03850256 -0.30345684 ...  0.14567922 -0.09989041
   -0.01025168]
 [ 0.18762272  0.10977384 -0.00291614 ... -0.39387344 -0.10644632
   -0.02397595]
 [ 0.10356236  0.0250188   0.09475959 ... -0.39387344 -0.05072119
   -0.01968712]
 ...
 [-0.13180663 -0.09826126 -0.0279612  ... -0.39387344 -0.09005657
   -0.02654925]
 [-0.13180663 -0.13486004  0.02212892 ...  0.14567922 -0.063833
   -0.02654925]
 [ 0.20443478  0.07895383 -0.17823155 ...  0.14567922 -0.09333451
   -0.02912255]]
```

Checking shap plots for LR1...

Expected value for LR: -0.6091565598361125
Summary Plot for SHAP Values in Test Set:

SHAP Bar Plot for SHAP Values Test Set:

SHAP Decision Plot for SHAP Values in Test Set:

SHAP Decision Plot for Single-Prediction in Test Set:

Checking feature importance for LR1...

Checking for list of csv files...

```
['/Users/jessicakim/Desktop/STREAMLINE/DemoData/Output/hcc_demo/hcc-data_example/model_evaluation/shap_values/testResu
lts/LR_1_shapFIValues_Test.csv']
```

Unnamed: 0	Features	Importance
0	Ascites degree*	0.242653
1	Symptoms	0.226906
2	Haemoglobin (g/dL)	0.223673
3	Alkaline phosphatase (U/L)	0.221572
4	Alpha-Fetoprotein (ng/mL)	0.216564
5	International Normalised Ratio*	0.202692
6	Performance Status*	0.195256
7	Age at diagnosis	0.166543
8	Diabetes	0.156453
9	Iron	0.155098
10	Albumin (mg/dL)	0.146038
11	Ferritin (ng/mL)	0.142575
12	Hepatitis C Virus Antibody	0.136120
13	Endemic Countries	0.121705
14	Mean Corpuscular Volume	0.112992
15	Smoking	0.109777
16	Major dimension of nodule (cm)	0.102912
17	Portal Vein Thrombosis	0.096549
18	Aspartate transaminase (U/L)	0.095388
19	Nonalcoholic Steatohepatitis	0.094967
20	Total Bilirubin(mg/dL)	0.083784
21	Portal Hypertension	0.077564
22	Direct Bilirubin (mg/dL)	0.077176
23	Leukocytes(G/L)	0.075128
24	Obesity	0.074326
25	Alanine transaminase (U/L)	0.072664
26	Packs of cigarets per year	0.056175
27	Gamma glutamyl transferase (U/L)	0.054443
28	Creatinine (mg/dL)	0.047948
29	Esophageal Varices	0.034705

30	30	Hepatitis B Core Antibody	0.029008
31	31	Total Proteins (g/dL)	0.025841
32	32	Platelets	0.022258
33	33	Cirrhosis	0.019556
34	34	Number of Nodules	0.017841
35	35	Grams of Alcohol per day	0.011571
36	36	Hepatitis B e Antigen	0.011481
37	37	Hepatitis B Surface Antigen	0.011007
38	38	Oxygen Saturation (%)	0.009104
39	39	Liver Metastasis	0.006661
40	40	Alcohol	0.004921

Out [18]:

Unnamed: 0		Features	Importance
0	0	Ascites degree*	0.242653
1	1	Symptoms	0.226906
2	2	Haemoglobin (g/dL)	0.223673
3	3	Alkaline phosphatase (U/L)	0.221572
4	4	Alpha-Fetoprotein (ng/mL)	0.216564
5	5	International Normalised Ratio*	0.202692
6	6	Performance Status*	0.195256
7	7	Age at diagnosis	0.166543
8	8	Diabetes	0.156453
9	9	Iron	0.155098
10	10	Albumin (mg/dL)	0.146038
11	11	Ferritin (ng/mL)	0.142575
12	12	Hepatitis C Virus Antibody	0.136120
13	13	Endemic Countries	0.121705
14	14	Mean Corpuscular Volume	0.112992
15	15	Smoking	0.109777
16	16	Major dimension of nodule (cm)	0.102912
17	17	Portal Vein Thrombosis	0.096549
18	18	Aspartate transaminase (U/L)	0.095388
19	19	Nonalcoholic Steatohepatitis	0.094967
20	20	Total Bilirubin(mg/dL)	0.083784
21	21	Portal Hypertension	0.077564
22	22	Direct Bilirubin (mg/dL)	0.077176
23	23	Leukocytes(G/L)	0.075128
24	24	Obesity	0.074326
25	25	Alanine transaminase (U/L)	0.072664
26	26	Packs of cigarets per year	0.056175
27	27	Gamma glutamyl transferase (U/L)	0.054443
28	28	Creatinine (mg/dL)	0.047948
29	29	Esophageal Varices	0.034705
30	30	Hepatitis B Core Antibody	0.029008
31	31	Total Proteins (g/dL)	0.025841
32	32	Platelets	0.022258
33	33	Cirrhosis	0.019556
34	34	Number of Nodules	0.017841
35	35	Grams of Alcohol per day	0.011571
36	36	Hepatitis B e Antigen	0.011481
37	37	Hepatitis B Surface Antigen	0.011007
38	38	Oxygen Saturation (%)	0.009104
39	39	Liver Metastasis	0.006661
40	40	Alcohol	0.004921

LR2 In CV2...

```
Checking if correct model is loaded...
LogisticRegression(C=0.0006580360277501316, class_weight='balanced', dual=True,
                    max_iter=112.07606211860569, random_state=42,
                    solver='liblinear')
['Age at diagnosis', 'Albumin (mg/dL)', 'Alcohol', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/mL)', 'Arteria
l Hypertension', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Chronic Renal Insufficiency', 'Creatinine (mg/d
L)', 'Diabetes', 'Direct Bilirubin (mg/dL)', 'Encephalopathy degree*', 'Endemic Countries', 'Ferritin (ng/mL)', 'Gamma
glutamyl transferase (U/L)', 'Haemoglobin (g/dL)', 'Hepatitis B Core Antibody', 'Hepatitis B e Antigen', 'Hepatitis C
Virus Antibody', 'Human Immunodeficiency Virus', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver
Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Number of Nodules', 'Obesity', 'Oxygen Satu
ration (%)', 'Packs of cigarets per year', 'Performance Status*', 'Platelets', 'Portal Hypertension', 'Portal Vein Thr
ombosis', 'Symptoms ']
['Age at diagnosis', 'Albumin (mg/dL)', 'Alcohol', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/mL)', 'Arteria
l Hypertension', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Chronic Renal Insufficiency', 'Creatinine (mg/d
L)', 'Diabetes', 'Direct Bilirubin (mg/dL)', 'Encephalopathy degree*', 'Endemic Countries', 'Ferritin (ng/mL)', 'Gamma
glutamyl transferase (U/L)', 'Haemoglobin (g/dL)', 'Hepatitis B Core Antibody', 'Hepatitis B e Antigen', 'Hepatitis C
Virus Antibody', 'Human Immunodeficiency Virus', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver
Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Number of Nodules', 'Obesity', 'Oxygen Satu
ration (%)', 'Packs of cigarets per year', 'Performance Status*', 'Platelets', 'Portal Hypertension', 'Portal Vein Thr
ombosis', 'Symptoms ']
```

Checking explainer for LR2...
<shap.explainers._linear.Linear object at 0x7f88a73ff100>

Checking shap values for LR2...

```
[[[-0.00209083  0.00438031  0.00118267 ... -0.00203892 -0.00439287
   -0.00869176]
 [-0.00982644  0.00438031  0.00118267 ...  0.00524293 -0.00439287
   0.00355015]
 [-0.00275388  0.00044561 -0.00473069 ... -0.00203892 -0.00439287
   0.00355015]
 ...
 [-0.00275388  0.01749597  0.00118267 ... -0.00203892 -0.00439287
   0.00355015]
 [-0.00319591 -0.00873536  0.00118267 ... -0.00203892 -0.00439287
   0.00355015]
 [-0.00010167  0.01880754  0.00118267 ... -0.00203892 -0.00439287
   0.00355015]]
```

Checking shap plots for LR2...

Expected value for LR: -0.006133751932115765
Summary Plot for SHAP Values in Test Set:

SHAP Bar Plot for SHAP Values Test Set:

SHAP Decision Plot for SHAP Values in Test Set:

SHAP Decision Plot for Single-Prediction in Test Set:

Checking feature importance for LR2...

Checking for list of csv files...

```
['/Users/jessicakim/Desktop/STREAMLINE/DemoData/Output/hcc_demo/hcc-data_example/model_evaluation/shap_values/testResu
lts/LR_2_shapFIValues_Test.csv']
```

Unnamed: 0	Features	Importance
0	Performance Status*	0.010779
1	Liver Metastasis	0.008865
2	Alkaline phosphatase (U/L)	0.007770
3	Ferritin (ng/mL)	0.007539
4	Portal Vein Thrombosis	0.007268
5	Albumin (mg/dL)	0.007223
6	Haemoglobin (g/dL)	0.006638
7	Ascites degree*	0.006479
8	Leukocytes(G/L)	0.006260
9	Number of Nodules	0.005690
10	Major dimension of nodule (cm)	0.005648
11	Iron	0.005371
12	Diabetes	0.005279
13	Symptoms	0.004765
14	Direct Bilirubin (mg/dL)	0.004723
15	Aspartate transaminase (U/L)	0.004438
16	Platelets	0.004057
17	Arterial Hypertension	0.003919
18	Hepatitis B Core Antibody	0.003809
19	Gamma glutamyl transferase (U/L)	0.003771
20	Obesity	0.003759
21	International Normalised Ratio*	0.003560
22	Portal Hypertension	0.002796
23	Human Immunodeficiency Virus	0.002731
24	Age at diagnosis	0.002526
25	Alcohol	0.002215
26	Mean Corpuscular Volume	0.002095
27	Hepatitis C Virus Antibody	0.001915
28	Chronic Renal Insufficiency	0.001647
29	Creatinine (mg/dL)	0.000991
30	Hepatitis B e Antigen	0.000957

31	31	Encephalopathy degree*	0.000757
32	32	Packs of cigarets per year	0.000481
33	33	Oxygen Saturation (%)	0.000419
34	34	Endemic Countries	0.000331
35	35	Alpha-Fetoprotein (ng/mL)	0.000059

Out [18]:

Unnamed: 0		Features	Importance
0	0	Performance Status*	0.010779
1	1	Liver Metastasis	0.008865
2	2	Alkaline phosphatase (U/L)	0.007770
3	3	Ferritin (ng/mL)	0.007539
4	4	Portal Vein Thrombosis	0.007268
5	5	Albumin (mg/dL)	0.007223
6	6	Haemoglobin (g/dL)	0.006638
7	7	Ascites degree*	0.006479
8	8	Leukocytes(G/L)	0.006260
9	9	Number of Nodules	0.005690
10	10	Major dimension of nodule (cm)	0.005648
11	11	Iron	0.005371
12	12	Diabetes	0.005279
13	13	Symptoms	0.004765
14	14	Direct Bilirubin (mg/dL)	0.004723
15	15	Aspartate transaminase (U/L)	0.004438
16	16	Platelets	0.004057
17	17	Arterial Hypertension	0.003919
18	18	Hepatitis B Core Antibody	0.003809
19	19	Gamma glutamyl transferase (U/L)	0.003771
20	20	Obesity	0.003759
21	21	International Normalised Ratio*	0.003560
22	22	Portal Hypertension	0.002796
23	23	Human Immunodeficiency Virus	0.002731
24	24	Age at diagnosis	0.002526
25	25	Alcohol	0.002215
26	26	Mean Corpuscular Volume	0.002095
27	27	Hepatitis C Virus Antibody	0.001915
28	28	Chronic Renal Insufficiency	0.001647
29	29	Creatinine (mg/dL)	0.000991
30	30	Hepatitis B e Antigen	0.000957
31	31	Encephalopathy degree*	0.000757
32	32	Packs of cigarets per year	0.000481
33	33	Oxygen Saturation (%)	0.000419
34	34	Endemic Countries	0.000331
35	35	Alpha-Fetoprotein (ng/mL)	0.000059

DT
DT0 In CV0...

Checking if correct model is loaded...
DecisionTreeClassifier(max_depth=17, min_samples_leaf=35, min_samples_split=45, random_state=42)
['Alanine transaminase (U/L)', 'Albumin (mg/dL)', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/mL)', 'Arterial Hypertension', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Chronic Renal Insufficiency', 'Cirrhosis', 'Creatinine (mg/dL)', 'Diabetes', 'Direct Bilirubin (mg/dL)', 'Encephalopathy degree*', 'Endemic Countries', 'Esophageal Varices', 'Ferritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haemoglobin (g/dL)', 'Hemochromatosis', 'Hepatitis B Surface Antigen', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibody', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Number of Nodules', 'Obesity', 'Oxygen Saturation (%)', 'Packs of cigarettes per year', 'Performance Status*', 'Portal Hypertension', 'Portal Vein Thrombosis', 'Smoking', 'Splenomegaly', 'Symptoms ', 'Total Bilirubin(mg/dL)']
['Alanine transaminase (U/L)', 'Albumin (mg/dL)', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/mL)', 'Arterial Hypertension', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Chronic Renal Insufficiency', 'Cirrhosis', 'Creatinine (mg/dL)', 'Diabetes', 'Direct Bilirubin (mg/dL)', 'Encephalopathy degree*', 'Endemic Countries', 'Esophageal Varices', 'Ferritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haemoglobin (g/dL)', 'Hemochromatosis', 'Hepatitis B Surface Antigen', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibody', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Number of Nodules', 'Obesity', 'Oxygen Saturation (%)', 'Packs of cigarettes per year', 'Performance Status*', 'Portal Hypertension', 'Portal Vein Thrombosis', 'Smoking', 'Splenomegaly', 'Symptoms ', 'Total Bilirubin(mg/dL)']

Checking explainer for DT0...
<shap.explainers._tree.Tree object at 0x7f88ca19da30>

Checking shap values for DT0...

```
[array([[0., 0., 0., ..., 0., 0., 0.],
        [0., 0., 0., ..., 0., 0., 0.],
        [0., 0., 0., ..., 0., 0., 0.],
        ...,
        [0., 0., 0., ..., 0., 0., 0.],
        [0., 0., 0., ..., 0., 0., 0.],
        [0., 0., 0., ..., 0., 0., 0.])), array([[0., 0., 0., ..., 0., 0., 0.],
        [0., 0., 0., ..., 0., 0., 0.],
        [0., 0., 0., ..., 0., 0., 0.],
        ...,
        [0., 0., 0., ..., 0., 0., 0.],
        [0., 0., 0., ..., 0., 0., 0.],
        [0., 0., 0., ..., 0., 0., 0.]])]
```

Checking shap plots for DT0...

Expected value for DT: [0.57272727 0.42727273]
Bar Summary Plot for SHAP Values in Class 0 & 1 in Test Set:

Decision Plot for SHAP Values from Class 0 in Test Set:

Decision Plot for SHAP Values from Class 1 in Test Set:

Checking feature importance for DT0...

Checking for list of csv files...

['/Users/jessicakim/Desktop/STREAMLINE/DemoData/Output/hcc_demo/hcc-data_example/model_evaluation/shap_values/testResults/DT_0_shapFIValues_Test.csv']

Unnamed: 0	Features	Importance
0	1 Alpha-Fetoprotein (ng/mL)	0.450314
1	2 Alanine transaminase (U/L)	0.000000
2	3 Obesity	0.000000
3	4 International Normalised Ratio*	0.000000
4	5 Iron	0.000000
5	6 Leukocytes(G/L)	0.000000
6	7 Liver Metastasis	0.000000
7	8 Major dimension of nodule (cm)	0.000000
8	9 Mean Corpuscular Volume	0.000000
9	10 Number of Nodules	0.000000
10	11 Oxygen Saturation (%)	0.000000
11	12 Hepatitis B e Antigen	0.000000
12	13 Packs of cigarettes per year	0.000000
13	14 Performance Status*	0.000000
14	15 Portal Hypertension	0.000000
15	16 Portal Vein Thrombosis	0.000000
16	17 Smoking	0.000000
17	18 Splenomegaly	0.000000
18	19 Symptoms	0.000000
19	20 Hepatitis C Virus Antibody	0.000000
20	21 Hepatitis B Surface Antigen	0.000000
21	22 Albumin (mg/dL)	0.000000
22	23 Hemochromatosis	0.000000
23	24 Alkaline phosphatase (U/L)	0.000000
24	25 Arterial Hypertension	0.000000
25	26 Ascites degree*	0.000000
26	27 Aspartate transaminase (U/L)	0.000000
27	28 Chronic Renal Insufficiency	0.000000
28	29 Cirrhosis	0.000000
29	30 Creatinine (mg/dL)	0.000000
30	31 Diabetes	0.000000

31	32	Direct Bilirubin (mg/dL)	0.000000
32	33	Encephalopathy degree*	0.000000
33	34	Endemic Countries	0.000000
34	35	Esophageal Varices	0.000000
35	36	Ferritin (ng/mL)	0.000000
36	37	Gamma glutamyl transferase (U/L)	0.000000
37	38	Haemoglobin (g/dL)	0.000000
38	39	Total Bilirubin(mg/dL)	0.000000

Out [18]:

Unnamed: 0		Features	Importance
0	1	Alpha-Fetoprotein (ng/mL)	0.450314
1	2	Alanine transaminase (U/L)	0.000000
2	3	Obesity	0.000000
3	4	International Normalised Ratio*	0.000000
4	5	Iron	0.000000
5	6	Leukocytes(G/L)	0.000000
6	7	Liver Metastasis	0.000000
7	8	Major dimension of nodule (cm)	0.000000
8	9	Mean Corpuscular Volume	0.000000
9	10	Number of Nodules	0.000000
10	11	Oxygen Saturation (%)	0.000000
11	12	Hepatitis B e Antigen	0.000000
12	13	Packs of cigarets per year	0.000000
13	14	Performance Status*	0.000000
14	15	Portal Hypertension	0.000000
15	16	Portal Vein Thrombosis	0.000000
16	17	Smoking	0.000000
17	18	Splenomegaly	0.000000
18	19	Symptoms	0.000000
19	20	Hepatitis C Virus Antibody	0.000000
20	21	Hepatitis B Surface Antigen	0.000000
21	22	Albumin (mg/dL)	0.000000
22	23	Hemochromatosis	0.000000
23	24	Alkaline phosphatase (U/L)	0.000000
24	25	Arterial Hypertension	0.000000
25	26	Ascites degree*	0.000000
26	27	Aspartate transaminase (U/L)	0.000000
27	28	Chronic Renal Insufficiency	0.000000
28	29	Cirrhosis	0.000000
29	30	Creatinine (mg/dL)	0.000000
30	31	Diabetes	0.000000
31	32	Direct Bilirubin (mg/dL)	0.000000
32	33	Encephalopathy degree*	0.000000
33	34	Endemic Countries	0.000000
34	35	Esophageal Varices	0.000000
35	36	Ferritin (ng/mL)	0.000000
36	37	Gamma glutamyl transferase (U/L)	0.000000
37	38	Haemoglobin (g/dL)	0.000000
38	39	Total Bilirubin(mg/dL)	0.000000

DT1 In CV1...

```
Checking if correct model is loaded...
DecisionTreeClassifier(criterion='entropy', max_depth=21, min_samples_leaf=3,
                        min_samples_split=23, random_state=42,
                        splitter='random')

['Age at diagnosis', 'Alanine transaminase (U/L)', 'Albumin (mg/dL)', 'Alcohol', 'Alkaline phosphatase (U/L)', 'Alpha-
Fetoprotein (ng/mL)', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Cirrhosis', 'Creatinine (mg/dL)', 'Diabete
s', 'Direct Bilirubin (mg/dL)', 'Endemic Countries', 'Esophageal Varices', 'Ferritin (ng/mL)', 'Gamma glutamyl transfe
rase (U/L)', 'Grams of Alcohol per day', 'Haemoglobin (g/dL)', 'Hepatitis B Core Antibody', 'Hepatitis B Surface Antig
en', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibody', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/
L)', 'Liver Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Nonalcoholic Steatohepatitis',
'Number of Nodules', 'Obesity', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Platele
ts', 'Portal Hypertension', 'Portal Vein Thrombosis', 'Smoking', 'Symptoms ', 'Total Bilirubin(mg/dL)', 'Total Protein
s (g/dL)']

['Age at diagnosis', 'Alanine transaminase (U/L)', 'Albumin (mg/dL)', 'Alcohol', 'Alkaline phosphatase (U/L)', 'Alpha-
Fetoprotein (ng/mL)', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Cirrhosis', 'Creatinine (mg/dL)', 'Diabete
s', 'Direct Bilirubin (mg/dL)', 'Endemic Countries', 'Esophageal Varices', 'Ferritin (ng/mL)', 'Gamma glutamyl transfe
rase (U/L)', 'Grams of Alcohol per day', 'Haemoglobin (g/dL)', 'Hepatitis B Core Antibody', 'Hepatitis B Surface Antig
en', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibody', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/
L)', 'Liver Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Nonalcoholic Steatohepatitis',
'Number of Nodules', 'Obesity', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Platele
ts', 'Portal Hypertension', 'Portal Vein Thrombosis', 'Smoking', 'Symptoms ', 'Total Bilirubin(mg/dL)', 'Total Protein
s (g/dL)']
```

Checking explainer for DT1...
<shap.explainers._tree.Tree object at 0x7f88ca989100>

Checking shap values for DT1...

```
[array([[ 0.          ,  0.          ,  0.          , ..., -0.02795699,
         0.          ,  0.          ],
       [ 0.          ,  0.          ,  0.          , ...,  0.07004662,
         0.          ,  0.          ],
       [ 0.          ,  0.          ,  0.          , ...,  0.07004662,
         0.          ,  0.          ],
       ...,
       [ 0.          ,  0.          ,  0.          , ...,  0.08666667,
         0.          ,  0.          ],
       [ 0.          ,  0.          ,  0.          , ..., -0.11345397,
         0.          ,  0.          ],
       [ 0.          ,  0.          ,  0.          , ..., -0.02795699,
         0.          ,  0.          ]]), array([[ 0.          ,  0.          ,  0.          , ...,  0.02795699,
         0.          ,  0.          ],
       [ 0.          ,  0.          ,  0.          , ..., -0.07004662,
         0.          ,  0.          ],
       [ 0.          ,  0.          ,  0.          , ..., -0.07004662,
         0.          ,  0.          ],
       ...,
       [ 0.          ,  0.          ,  0.          , ..., -0.08666667,
         0.          ,  0.          ],
       [ 0.          ,  0.          ,  0.          , ...,  0.11345397,
         0.          ,  0.          ],
       [ 0.          ,  0.          ,  0.          , ...,  0.02795699,
         0.          ,  0.          ]]])
```

Checking shap plots for DT1...

Expected value for DT: [0.63636364 0.36363636]
Bar Summary Plot for SHAP Values in Class 0 & 1 in Test Set:

Decision Plot for SHAP Values from Class 0 in Test Set:

Decision Plot for SHAP Values from Class 1 in Test Set:

Checking feature importance for DT1...

Checking for list of csv files...

```
['/Users/jessicakim/Desktop/STREAMLINE/DemoData/Output/hcc_demo/hcc-data_example/model_evaluation/shap_values/testResu
lts/DT_1_shapFIValues_Test.csv']
```

Unnamed: 0	Features	Importance
0	1 Ascites degree*	0.307172
1	2 International Normalised Ratio*	0.250056
2	3 Performance Status*	0.156491
3	4 Alcohol	0.135875
4	5 Symptoms	0.127666
5	6 Haemoglobin (g/dL)	0.089932
6	7 Mean Corpuscular Volume	0.055060
7	8 Gamma glutamyl transferase (U/L)	0.046579
8	9 Alpha-Fetoprotein (ng/mL)	0.040349
9	10 Obesity	0.000000
10	11 Liver Metastasis	0.000000
11	12 Major dimension of nodule (cm)	0.000000
12	13 Nonalcoholic Steatohepatitis	0.000000
13	14 Number of Nodules	0.000000
14	15 Oxygen Saturation (%)	0.000000
15	16 Packs of cigarets per year	0.000000
16	17 Leukocytes(G/L)	0.000000

17	18	Platelets	0.000000
18	19	Portal Hypertension	0.000000
19	20	Portal Vein Thrombosis	0.000000
20	21	Smoking	0.000000
21	22	Total Bilirubin(mg/dL)	0.000000
22	23	Age at diagnosis	0.000000
23	24	Hepatitis B e Antigen	0.000000
24	25	Iron	0.000000
25	26	Direct Bilirubin (mg/dL)	0.000000
26	27	Albumin (mg/dL)	0.000000
27	28	Alkaline phosphatase (U/L)	0.000000
28	29	Aspartate transaminase (U/L)	0.000000
29	30	Cirrhosis	0.000000
30	31	Creatinine (mg/dL)	0.000000
31	32	Diabetes	0.000000
32	33	Endemic Countries	0.000000
33	34	Hepatitis C Virus Antibody	0.000000
34	35	Esophageal Varices	0.000000
35	36	Ferritin (ng/mL)	0.000000
36	37	Grams of Alcohol per day	0.000000
37	38	Hepatitis B Core Antibody	0.000000
38	39	Hepatitis B Surface Antigen	0.000000
39	40	Alanine transaminase (U/L)	0.000000
40	41	Total Proteins (g/dL)	0.000000

Out [18] :

Unnamed: 0		Features	Importance
0	1	Ascites degree*	0.307172
1	2	International Normalised Ratio*	0.250056
2	3	Performance Status*	0.156491
3	4	Alcohol	0.135875
4	5	Symptoms	0.127666
5	6	Haemoglobin (g/dL)	0.089932
6	7	Mean Corpuscular Volume	0.055060
7	8	Gamma glutamyl transferase (U/L)	0.046579
8	9	Alpha-Fetoprotein (ng/mL)	0.040349
9	10	Obesity	0.000000
10	11	Liver Metastasis	0.000000
11	12	Major dimension of nodule (cm)	0.000000
12	13	Nonalcoholic Steatohepatitis	0.000000
13	14	Number of Nodules	0.000000
14	15	Oxygen Saturation (%)	0.000000
15	16	Packs of cigarets per year	0.000000
16	17	Leukocytes(G/L)	0.000000
17	18	Platelets	0.000000
18	19	Portal Hypertension	0.000000
19	20	Portal Vein Thrombosis	0.000000
20	21	Smoking	0.000000
21	22	Total Bilirubin(mg/dL)	0.000000
22	23	Age at diagnosis	0.000000
23	24	Hepatitis B e Antigen	0.000000
24	25	Iron	0.000000
25	26	Direct Bilirubin (mg/dL)	0.000000
26	27	Albumin (mg/dL)	0.000000
27	28	Alkaline phosphatase (U/L)	0.000000
28	29	Aspartate transaminase (U/L)	0.000000
29	30	Cirrhosis	0.000000
30	31	Creatinine (mg/dL)	0.000000
31	32	Diabetes	0.000000
32	33	Endemic Countries	0.000000
33	34	Hepatitis C Virus Antibody	0.000000
34	35	Esophageal Varices	0.000000
35	36	Ferritin (ng/mL)	0.000000
36	37	Grams of Alcohol per day	0.000000
37	38	Hepatitis B Core Antibody	0.000000
38	39	Hepatitis B Surface Antigen	0.000000
39	40	Alanine transaminase (U/L)	0.000000
40	41	Total Proteins (g/dL)	0.000000

DT2 In CV2...

```
Checking if correct model is loaded...
DecisionTreeClassifier(class_weight='balanced', max_depth=29,
                        min_samples_leaf=30, min_samples_split=45,
                        random_state=42)
['Age at diagnosis', 'Albumin (mg/dL)', 'Alcohol', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/mL)', 'Arterial Hypertension', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Chronic Renal Insufficiency', 'Creatinine (mg/dL)', 'Diabetes', 'Direct Bilirubin (mg/dL)', 'Encephalopathy degree*', 'Endemic Countries', 'Ferritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haemoglobin (g/dL)', 'Hepatitis B Core Antibody', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibody', 'Human Immunodeficiency Virus', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Number of Nodules', 'Obesity', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Platelets', 'Portal Hypertension', 'Portal Vein Thrombosis', 'Symptoms ']
```

Checking explainer for DT2...
<shap.explainers._tree.Tree object at 0x7f88ca97fbb0>

Checking shap values for DT2...

```
[array([[ 0.          , -0.14385676,  0.          , ...,  0.          ,
         0.          ,  0.          ],
       [ 0.          , -0.0462963 ,  0.          , ...,  0.          ,
         0.          ,  0.          ],
       [ 0.          , -0.0462963 ,  0.          , ...,  0.          ,
         0.          ,  0.          ],
       ...,
       [ 0.          , -0.0462963 ,  0.          , ...,  0.          ,
         0.          ,  0.          ],
       [ 0.          ,  0.05769231,  0.          , ...,  0.          ,
         0.          ,  0.          ],
       [ 0.          , -0.14385676,  0.          , ...,  0.          ,
         0.          ,  0.          ]]), array([[ 0.          ,  0.14385676,  0.          , ...,  0.          ,
         0.          ,  0.          ],
       [ 0.          ,  0.0462963 ,  0.          , ...,  0.          ,
         0.          ,  0.          ],
       [ 0.          ,  0.0462963 ,  0.          , ...,  0.          ,
         0.          ,  0.          ],
       ...,
       [ 0.          ,  0.0462963 ,  0.          , ...,  0.          ,
         0.          ,  0.          ],
       [ 0.          , -0.05769231,  0.          , ...,  0.          ,
         0.          ,  0.          ],
       [ 0.          ,  0.14385676,  0.          , ...,  0.          ,
         0.          ,  0.          ]]])]
```

Checking shap plots for DT2...

Expected value for DT: [0.5 0.5]
Bar Summary Plot for SHAP Values in Class 0 & 1 in Test Set:

Decision Plot for SHAP Values from Class 0 in Test Set:

Decision Plot for SHAP Values from Class 1 in Test Set:

Checking feature importance for DT2...

Checking for list of csv files...

```
['/Users/jessicakim/Desktop/STREAMLINE/DemoData/Output/hcc_demo/hcc-data_example/model_evaluation/shap_values/testResults/DT_2_shapFIValues_Test.csv']
```

Unnamed: 0		Features	Importance
0	1	Alkaline phosphatase (U/L)	0.455851
1	2	Albumin (mg/dL)	0.204616
2	3	Number of Nodules	0.000000
3	4	Human Immunodeficiency Virus	0.000000
4	5	International Normalised Ratio*	0.000000
5	6	Iron	0.000000
6	7	Leukocytes(G/L)	0.000000
7	8	Liver Metastasis	0.000000
8	9	Major dimension of nodule (cm)	0.000000
9	10	Mean Corpuscular Volume	0.000000
10	11	Age at diagnosis	0.000000
11	12	Hepatitis C Virus Antibody	0.000000
12	13	Oxygen Saturation (%)	0.000000
13	14	Packs of cigarets per year	0.000000
14	15	Performance Status*	0.000000
15	16	Platelets	0.000000
16	17	Portal Hypertension	0.000000
17	18	Portal Vein Thrombosis	0.000000
18	19	Obesity	0.000000

19	20	Hepatitis B e Antigen	0.000000
20	21	Hepatitis B Core Antibody	0.000000
21	22	Haemoglobin (g/dL)	0.000000
22	23	Gamma glutamyl transferase (U/L)	0.000000
23	24	Ferritin (ng/mL)	0.000000
24	25	Endemic Countries	0.000000
25	26	Encephalopathy degree*	0.000000
26	27	Direct Bilirubin (mg/dL)	0.000000
27	28	Diabetes	0.000000
28	29	Creatinine (mg/dL)	0.000000
29	30	Chronic Renal Insufficiency	0.000000
30	31	Aspartate transaminase (U/L)	0.000000
31	32	Ascites degree*	0.000000
32	33	Arterial Hypertension	0.000000
33	34	Alpha-Fetoprotein (ng/mL)	0.000000
34	35	Alcohol	0.000000
35	36	Symptoms	0.000000

Out [18]:

Unnamed: 0		Features	Importance
0	1	Alkaline phosphatase (U/L)	0.455851
1	2	Albumin (mg/dL)	0.204616
2	3	Number of Nodules	0.000000
3	4	Human Immunodeficiency Virus	0.000000
4	5	International Normalised Ratio*	0.000000
5	6	Iron	0.000000
6	7	Leukocytes(G/L)	0.000000
7	8	Liver Metastasis	0.000000
8	9	Major dimension of nodule (cm)	0.000000
9	10	Mean Corpuscular Volume	0.000000
10	11	Age at diagnosis	0.000000
11	12	Hepatitis C Virus Antibody	0.000000
12	13	Oxygen Saturation (%)	0.000000
13	14	Packs of cigarets per year	0.000000
14	15	Performance Status*	0.000000
15	16	Platelets	0.000000
16	17	Portal Hypertension	0.000000
17	18	Portal Vein Thrombosis	0.000000
18	19	Obesity	0.000000
19	20	Hepatitis B e Antigen	0.000000
20	21	Hepatitis B Core Antibody	0.000000
21	22	Haemoglobin (g/dL)	0.000000
22	23	Gamma glutamyl transferase (U/L)	0.000000
23	24	Ferritin (ng/mL)	0.000000
24	25	Endemic Countries	0.000000
25	26	Encephalopathy degree*	0.000000
26	27	Direct Bilirubin (mg/dL)	0.000000
27	28	Diabetes	0.000000
28	29	Creatinine (mg/dL)	0.000000
29	30	Chronic Renal Insufficiency	0.000000
30	31	Aspartate transaminase (U/L)	0.000000
31	32	Ascites degree*	0.000000
32	33	Arterial Hypertension	0.000000
33	34	Alpha-Fetoprotein (ng/mL)	0.000000
34	35	Alcohol	0.000000
35	36	Symptoms	0.000000

```
RF
RF0 In CV0...

Checking if correct model is loaded...
RandomForestClassifier(criterion='entropy', max_depth=1, max_features=None,
                        min_samples_leaf=17, min_samples_split=41,
                        n_estimators=960, random_state=42)
['Alanine transaminase (U/L)', 'Albumin (mg/dL)', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/mL)', 'Arterial Hypertension', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Chronic Renal Insufficiency', 'Cirrhosis', 'Creatinine (mg/dL)', 'Diabetes', 'Direct Bilirubin (mg/dL)', 'Encephalopathy degree*', 'Endemic Countries', 'Esophageal Varices', 'Ferritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haemoglobin (g/dL)', 'Hemochromatosis', 'Hepatitis B Surface Antigen', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibody', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Number of Nodules', 'Obesity', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Portal Hypertension', 'Portal Vein Thrombosis', 'Smoking', 'Splenomegaly', 'Symptoms ', 'Total Bilirubin(mg/dL)']
['Alanine transaminase (U/L)', 'Albumin (mg/dL)', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/mL)', 'Arterial Hypertension', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Chronic Renal Insufficiency', 'Cirrhosis', 'Creatinine (mg/dL)', 'Diabetes', 'Direct Bilirubin (mg/dL)', 'Encephalopathy degree*', 'Endemic Countries', 'Esophageal Varices', 'Ferritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haemoglobin (g/dL)', 'Hemochromatosis', 'Hepatitis B Surface Antigen', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibody', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Number of Nodules', 'Obesity', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Portal Hypertension', 'Portal Vein Thrombosis', 'Smoking', 'Splenomegaly', 'Symptoms ', 'Total Bilirubin(mg/dL)']
```

Checking explainer for RF0...
<shap.explainers._tree.Tree object at 0x7f88ca13bb50>

Checking shap values for RF0...

```
[array([[ 0.          ,  0.04181896,  0.00365806, ...,  0.          ,
         0.          ,  0.          ],
        [ 0.          , -0.01898534, -0.01999966, ...,  0.          ,
         0.          ,  0.          ],
        [ 0.          , -0.01898534, -0.01840236, ...,  0.          ,
         0.          ,  0.          ],
        ...,
        [ 0.          ,  0.04012049,  0.02614831, ...,  0.          ,
         0.          ,  0.          ],
        [ 0.          , -0.01898534, -0.00631506, ...,  0.          ,
         0.          ,  0.          ],
        [ 0.          , -0.01898534, -0.01061794, ...,  0.          ,
         0.          ,  0.          ]]), array([[ 0.          , -0.04181896, -0.00365806, ...,  0.          ,
         0.          ,  0.          ],
        [ 0.          ,  0.01898534,  0.01999966, ...,  0.          ,
         0.          ,  0.          ],
        [ 0.          ,  0.01898534,  0.01840236, ...,  0.          ,
         0.          ,  0.          ],
        ...,
        [ 0.          , -0.04012049, -0.02614831, ...,  0.          ,
         0.          ,  0.          ],
        [ 0.          ,  0.01898534,  0.00631506, ...,  0.          ,
         0.          ,  0.          ],
        [ 0.          ,  0.01898534,  0.01061794, ...,  0.          ,
         0.          ,  0.          ]]])
```

Checking shap plots for RF0...

Expected value for RF: [0.5728125 0.4271875]
Bar Summary Plot for SHAP Values in Class 0 & 1 in Test Set:

Decision Plot for SHAP Values from Class 0 in Test Set:

Decision Plot for SHAP Values from Class 1 in Test Set:

Checking feature importance for RF0...

Checking for list of csv files...

```
['/Users/jessicakim/Desktop/STREAMLINE/DemoData/Output/hcc_demo/hcc-data_example/model_evaluation/shap_values/testResults/RF_0_shapFIValues_Test.csv']
```

Unnamed: 0		Features	Importance
0	1	Alpha-Fetoprotein (ng/mL)	0.263811
1	2	Albumin (mg/dL)	0.057781
2	3	Alkaline phosphatase (U/L)	0.033483
3	4	Iron	0.022969
4	5	Performance Status*	0.010778
5	6	Aspartate transaminase (U/L)	0.007389
6	7	Ferritin (ng/mL)	0.005711
7	8	Haemoglobin (g/dL)	0.003896
8	9	Creatinine (mg/dL)	0.003621
9	10	Liver Metastasis	0.003396
10	11	Gamma glutamyl transferase (U/L)	0.003386
11	12	Direct Bilirubin (mg/dL)	0.002852
12	13	Major dimension of nodule (cm)	0.002410
13	14	Leukocytes(G/L)	0.001034
14	15	Oxygen Saturation (%)	0.000000
15	16	Packs of cigarets per year	0.000000
16	17	Portal Hypertension	0.000000
17	18	Portal Vein Thrombosis	0.000000

18	19	Obesity	0.000000
19	20	Smoking	0.000000
20	21	Number of Nodules	0.000000
21	22	Mean Corpuscular Volume	0.000000
22	23	Splenomegaly	0.000000
23	24	Symptoms	0.000000
24	25	Alanine transaminase (U/L)	0.000000
25	26	Hepatitis B Surface Antigen	0.000000
26	27	International Normalised Ratio*	0.000000
27	28	Hepatitis C Virus Antibody	0.000000
28	29	Hepatitis B e Antigen	0.000000
29	30	Hemochromatosis	0.000000
30	31	Esophageal Varices	0.000000
31	32	Endemic Countries	0.000000
32	33	Encephalopathy degree*	0.000000
33	34	Diabetes	0.000000
34	35	Cirrhosis	0.000000
35	36	Chronic Renal Insufficiency	0.000000
36	37	Ascites degree*	0.000000
37	38	Arterial Hypertension	0.000000
38	39	Total Bilirubin(mg/dL)	0.000000

Out [18]:

Unnamed: 0		Features	Importance
0	1	Alpha-Fetoprotein (ng/mL)	0.263811
1	2	Albumin (mg/dL)	0.057781
2	3	Alkaline phosphatase (U/L)	0.033483
3	4	Iron	0.022969
4	5	Performance Status*	0.010778
5	6	Aspartate transaminase (U/L)	0.007389
6	7	Ferritin (ng/mL)	0.005711
7	8	Haemoglobin (g/dL)	0.003896
8	9	Creatinine (mg/dL)	0.003621
9	10	Liver Metastasis	0.003396
10	11	Gamma glutamyl transferase (U/L)	0.003386
11	12	Direct Bilirubin (mg/dL)	0.002852
12	13	Major dimension of nodule (cm)	0.002410
13	14	Leukocytes(G/L)	0.001034
14	15	Oxygen Saturation (%)	0.000000
15	16	Packs of cigarets per year	0.000000
16	17	Portal Hypertension	0.000000
17	18	Portal Vein Thrombosis	0.000000
18	19	Obesity	0.000000
19	20	Smoking	0.000000
20	21	Number of Nodules	0.000000
21	22	Mean Corpuscular Volume	0.000000
22	23	Splenomegaly	0.000000
23	24	Symptoms	0.000000
24	25	Alanine transaminase (U/L)	0.000000
25	26	Hepatitis B Surface Antigen	0.000000
26	27	International Normalised Ratio*	0.000000
27	28	Hepatitis C Virus Antibody	0.000000
28	29	Hepatitis B e Antigen	0.000000
29	30	Hemochromatosis	0.000000
30	31	Esophageal Varices	0.000000
31	32	Endemic Countries	0.000000
32	33	Encephalopathy degree*	0.000000
33	34	Diabetes	0.000000
34	35	Cirrhosis	0.000000
35	36	Chronic Renal Insufficiency	0.000000
36	37	Ascites degree*	0.000000
37	38	Arterial Hypertension	0.000000
38	39	Total Bilirubin(mg/dL)	0.000000

RF1 In CV1...

```
Checking if correct model is loaded...
RandomForestClassifier(class_weight='balanced', criterion='entropy',
                        max_depth=2, max_features='log2', min_samples_leaf=9,
                        min_samples_split=31, n_estimators=207, random_state=42)
['Age at diagnosis', 'Alanine transaminase (U/L)', 'Albumin (mg/dL)', 'Alcohol', 'Alkaline phosphatase (U/L)', 'Alpha-
Fetoprotein (ng/mL)', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Cirrhosis', 'Creatinine (mg/dL)', 'Diabete
s', 'Direct Bilirubin (mg/dL)', 'Endemic Countries', 'Esophageal Varices', 'Ferritin (ng/mL)', 'Gamma glutamyl transfe
rase (U/L)', 'Grams of Alcohol per day', 'Haemoglobin (g/dL)', 'Hepatitis B Core Antibody', 'Hepatitis B Surface Antig
en', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibody', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/
L)', 'Liver Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Nonalcoholic Steatohepatitis',
'Number of Nodules', 'Obesity', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Platele
ts', 'Portal Hypertension', 'Portal Vein Thrombosis', 'Smoking', 'Symptoms ', 'Total Bilirubin(mg/dL)', 'Total Protein
s (g/dL)']
['Age at diagnosis', 'Alanine transaminase (U/L)', 'Albumin (mg/dL)', 'Alcohol', 'Alkaline phosphatase (U/L)', 'Alpha-
Fetoprotein (ng/mL)', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Cirrhosis', 'Creatinine (mg/dL)', 'Diabete
s', 'Direct Bilirubin (mg/dL)', 'Endemic Countries', 'Esophageal Varices', 'Ferritin (ng/mL)', 'Gamma glutamyl transfe
rase (U/L)', 'Grams of Alcohol per day', 'Haemoglobin (g/dL)', 'Hepatitis B Core Antibody', 'Hepatitis B Surface Antig
en', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibody', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/
L)', 'Liver Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Nonalcoholic Steatohepatitis',
'Number of Nodules', 'Obesity', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Platele
ts', 'Portal Hypertension', 'Portal Vein Thrombosis', 'Smoking', 'Symptoms ', 'Total Bilirubin(mg/dL)', 'Total Protein
s (g/dL)']
```

Checking explainer for RF1...
<shap.explainers._tree.Tree object at 0x7f8907c99a90>

Checking shap values for RF1...

```
[array([[ 3.95998286e-03,  8.78166228e-05,  5.42231605e-02, ...,
        -4.99207302e-03,  7.28955367e-03,  2.80322166e-03],
        [-3.19678857e-03,  8.79064012e-04,  1.60776918e-03, ...,
         1.80742612e-02,  5.37725417e-03, -1.63881455e-03],
        [-2.91886671e-03, -4.37616095e-04, -2.66173097e-02, ...,
         2.10142415e-02,  3.35299303e-03,  5.93880632e-04],
        ...,
         [ 2.62833254e-03, -1.78372567e-04,  1.24812736e-02, ...,
          1.73079030e-02,  4.94635866e-03, -1.53940799e-03],
         [ 2.33387141e-03, -4.42763000e-04, -2.58700195e-02, ...,
          -6.86380765e-03,  4.07652440e-03, -1.52713434e-03],
         [-5.01472471e-03,  1.13411715e-03,  5.63221558e-02, ...,
          -5.51702530e-03,  5.22884231e-03, -4.18441541e-04]])], array([[ -3.95998286e-03, -8.78166228e-05, -5.42231605e-0
2, ...,
         4.99207302e-03, -7.28955367e-03, -2.80322166e-03],
         [ 3.19678857e-03, -8.79064012e-04, -1.60776918e-03, ...,
          -1.80742612e-02, -5.37725417e-03,  1.63881455e-03],
         [ 2.91886671e-03,  4.37616095e-04,  2.66173097e-02, ...,
          -2.10142415e-02, -3.35299303e-03, -5.93880632e-04],
         ...,
         [-2.62833254e-03,  1.78372567e-04, -1.24812736e-02, ...,
          -1.73079030e-02, -4.94635866e-03,  1.53940799e-03],
         [-2.33387141e-03,  4.42763000e-04,  2.58700195e-02, ...,
          6.86380765e-03, -4.07652440e-03,  1.52713434e-03],
         [ 5.01472471e-03, -1.13411715e-03, -5.63221558e-02, ...,
          5.51702530e-03, -5.22884231e-03,  4.18441541e-04]])])]
```

Checking shap plots for RF1...

Expected value for RF: [0.49673858 0.50326142]
Bar Summary Plot for SHAP Values in Class 0 & 1 in Test Set:

Decision Plot for SHAP Values from Class 0 in Test Set:

Decision Plot for SHAP Values from Class 1 in Test Set:

Checking feature importance for RF1...

Checking for list of csv files...

```
[ '/Users/jessicakim/Desktop/STREAMLINE/DemoData/Output/hcc_demo/hcc-data_example/model_evaluation/shap_values/testResults/RF_1_shapFIValues_Test.csv' ]
```

Unnamed: 0	Features	Importance	
0	1	Haemoglobin (g/dL)	0.065451
1	2	Albumin (mg/dL)	0.060464
2	3	Ascites degree*	0.055906
3	4	Iron	0.051997
4	5	Oxygen Saturation (%)	0.044540
5	6	Performance Status*	0.041592
6	7	Direct Bilirubin (mg/dL)	0.040724
7	8	Alkaline phosphatase (U/L)	0.034861
8	9	Alpha-Fetoprotein (ng/mL)	0.034022
9	10	International Normalised Ratio*	0.024608
10	11	Ferritin (ng/mL)	0.019971
11	12	Symptoms	0.019735
12	13	Mean Corpuscular Volume	0.014224
13	14	Total Bilirubin(mg/dL)	0.011621
14	15	Major dimension of nodule (cm)	0.010869
15	16	Packs of cigarets per year	0.007936

16	17	Aspartate transaminase (U/L)	0.007492
17	18	Age at diagnosis	0.006167
18	19	Creatinine (mg/dL)	0.005623
19	20	Gamma glutamyl transferase (U/L)	0.005091
20	21	Grams of Alcohol per day	0.003893
21	22	Platelets	0.003571
22	23	Total Proteins (g/dL)	0.003384
23	24	Leukocytes(G/L)	0.002181
24	25	Number of Nodules	0.001991
25	26	Smoking	0.001508
26	27	Diabetes	0.001237
27	28	Alanine transaminase (U/L)	0.001119
28	29	Portal Hypertension	0.001070
29	30	Hepatitis B Core Antibody	0.000886
30	31	Alcohol	0.000775
31	32	Hepatitis C Virus Antibody	0.000764
32	33	Portal Vein Thrombosis	0.000237
33	34	Cirrhosis	0.000000
34	35	Esophageal Varices	0.000000
35	36	Endemic Countries	0.000000
36	37	Obesity	0.000000
37	38	Nonalcoholic Steatohepatitis	0.000000
38	39	Hepatitis B Surface Antigen	0.000000
39	40	Liver Metastasis	0.000000
40	41	Hepatitis B e Antigen	0.000000

Out [18] :

Unnamed: 0		Features	Importance
0	1	Haemoglobin (g/dL)	0.065451
1	2	Albumin (mg/dL)	0.060464
2	3	Ascites degree*	0.055906
3	4	Iron	0.051997
4	5	Oxygen Saturation (%)	0.044540
5	6	Performance Status*	0.041592
6	7	Direct Bilirubin (mg/dL)	0.040724
7	8	Alkaline phosphatase (U/L)	0.034861
8	9	Alpha-Fetoprotein (ng/mL)	0.034022
9	10	International Normalised Ratio*	0.024608
10	11	Ferritin (ng/mL)	0.019971
11	12	Symptoms	0.019735
12	13	Mean Corpuscular Volume	0.014224
13	14	Total Bilirubin(mg/dL)	0.011621
14	15	Major dimension of nodule (cm)	0.010869
15	16	Packs of cigarets per year	0.007936
16	17	Aspartate transaminase (U/L)	0.007492
17	18	Age at diagnosis	0.006167
18	19	Creatinine (mg/dL)	0.005623
19	20	Gamma glutamyl transferase (U/L)	0.005091
20	21	Grams of Alcohol per day	0.003893
21	22	Platelets	0.003571
22	23	Total Proteins (g/dL)	0.003384
23	24	Leukocytes(G/L)	0.002181
24	25	Number of Nodules	0.001991
25	26	Smoking	0.001508
26	27	Diabetes	0.001237
27	28	Alanine transaminase (U/L)	0.001119
28	29	Portal Hypertension	0.001070
29	30	Hepatitis B Core Antibody	0.000886
30	31	Alcohol	0.000775
31	32	Hepatitis C Virus Antibody	0.000764
32	33	Portal Vein Thrombosis	0.000237
33	34	Cirrhosis	0.000000
34	35	Esophageal Varices	0.000000
35	36	Endemic Countries	0.000000
36	37	Obesity	0.000000
37	38	Nonalcoholic Steatohepatitis	0.000000
38	39	Hepatitis B Surface Antigen	0.000000
39	40	Liver Metastasis	0.000000
40	41	Hepatitis B e Antigen	0.000000

RF2 In CV2...

```
Checking if correct model is loaded...
RandomForestClassifier(max_depth=11, max_features=None, min_samples_leaf=14,
                        min_samples_split=27, n_estimators=10, random_state=42)
['Age at diagnosis', 'Albumin (mg/dL)', 'Alcohol', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/mL)', 'Arterial Hypertension', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Chronic Renal Insufficiency', 'Creatinine (mg/dL)', 'Diabetes', 'Direct Bilirubin (mg/dL)', 'Encephalopathy degree*', 'Endemic Countries', 'Ferritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haemoglobin (g/dL)', 'Hepatitis B Core Antibody', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibody', 'Human Immunodeficiency Virus', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Number of Nodules', 'Obesity', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Platelets', 'Portal Hypertension', 'Portal Vein Thrombosis', 'Symptoms ']
```

Checking explainer for RF2...
<shap.explainers._tree.Tree object at 0x7f88a5e17640>

Checking shap values for RF2...

```
[array([[ 0.01794027, -0.01378879,  0.          , ...,  0.          ,
         0.          ,  0.          ],
       [ 0.00711062, -0.00448347,  0.          , ...,  0.          ,
         0.          ,  0.          ],
       [ 0.00711062,  0.00318182,  0.          , ...,  0.          ,
         0.          ,  0.          ],
       ...,
       [ 0.00711062, -0.00448347,  0.          , ...,  0.          ,
         0.          ,  0.          ],
       [ 0.01210694,  0.00318182,  0.          , ...,  0.          ,
         0.          ,  0.          ],
       [ 0.01794027, -0.01378879,  0.          , ...,  0.          ,
         0.          ,  0.          ]]), array([[ -0.01794027,  0.01378879,  0.          , ...,  0.          ,
        -0.00711062,  0.00448347,  0.          , ...,  0.          ,
        -0.00711062, -0.00318182,  0.          , ...,  0.          ,
        -0.00711062,  0.00448347,  0.          , ...,  0.          ,
        -0.01210694, -0.00318182,  0.          , ...,  0.          ,
        -0.01794027,  0.01378879,  0.          , ...,  0.          ,
        0.          ,  0.          ]])]
```

Checking shap plots for RF2...

Expected value for RF: [0.61909091 0.38090909]
Bar Summary Plot for SHAP Values in Class 0 & 1 in Test Set:

Decision Plot for SHAP Values from Class 0 in Test Set:

Decision Plot for SHAP Values from Class 1 in Test Set:

Checking feature importance for RF2...

Checking for list of csv files...

```
['/Users/jessicakim/Desktop/STREAMLINE/DemoData/Output/hcc_demo/hcc-data_example/model_evaluation/shap_values/testResults/RF_2_shapFIValues_Test.csv']
```

Unnamed: 0		Features	Importance
0	1	Alkaline phosphatase (U/L)	0.343480
1	2	Ferritin (ng/mL)	0.119536
2	3	Iron	0.043435
3	4	Major dimension of nodule (cm)	0.040151
4	5	Number of Nodules	0.039206
5	6	Alpha-Fetoprotein (ng/mL)	0.035987
6	7	Age at diagnosis	0.029558
7	8	Ascites degree*	0.024177
8	9	Aspartate transaminase (U/L)	0.021923
9	10	Mean Corpuscular Volume	0.017171
10	11	Albumin (mg/dL)	0.015785
11	12	Performance Status*	0.015073
12	13	Packs of cigarets per year	0.008147
13	14	Gamma glutamyl transferase (U/L)	0.002236
14	15	Obesity	0.000000
15	16	Oxygen Saturation (%)	0.000000
16	17	Platelets	0.000000
17	18	Portal Hypertension	0.000000
18	19	Portal Vein Thrombosis	0.000000
19	20	Liver Metastasis	0.000000

20	21	Leukocytes(G/L)	0.000000
21	22	Hepatitis B e Antigen	0.000000
22	23	International Normalised Ratio*	0.000000
23	24	Human Immunodeficiency Virus	0.000000
24	25	Hepatitis C Virus Antibody	0.000000
25	26	Hepatitis B Core Antibody	0.000000
26	27	Haemoglobin (g/dL)	0.000000
27	28	Endemic Countries	0.000000
28	29	Encephalopathy degree*	0.000000
29	30	Direct Bilirubin (mg/dL)	0.000000
30	31	Diabetes	0.000000
31	32	Creatinine (mg/dL)	0.000000
32	33	Chronic Renal Insufficiency	0.000000
33	34	Arterial Hypertension	0.000000
34	35	Alcohol	0.000000
35	36	Symptoms	0.000000

Out [18]:

Unnamed: 0		Features	Importance
0	1	Alkaline phosphatase (U/L)	0.343480
1	2	Ferritin (ng/mL)	0.119536
2	3	Iron	0.043435
3	4	Major dimension of nodule (cm)	0.040151
4	5	Number of Nodules	0.039206
5	6	Alpha-Fetoprotein (ng/mL)	0.035987
6	7	Age at diagnosis	0.029558
7	8	Ascites degree*	0.024177
8	9	Aspartate transaminase (U/L)	0.021923
9	10	Mean Corpuscular Volume	0.017171
10	11	Albumin (mg/dL)	0.015785
11	12	Performance Status*	0.015073
12	13	Packs of cigarets per year	0.008147
13	14	Gamma glutamyl transferase (U/L)	0.002236
14	15	Obesity	0.000000
15	16	Oxygen Saturation (%)	0.000000
16	17	Platelets	0.000000
17	18	Portal Hypertension	0.000000
18	19	Portal Vein Thrombosis	0.000000
19	20	Liver Metastasis	0.000000
20	21	Leukocytes(G/L)	0.000000
21	22	Hepatitis B e Antigen	0.000000
22	23	International Normalised Ratio*	0.000000
23	24	Human Immunodeficiency Virus	0.000000
24	25	Hepatitis C Virus Antibody	0.000000
25	26	Hepatitis B Core Antibody	0.000000
26	27	Haemoglobin (g/dL)	0.000000
27	28	Endemic Countries	0.000000
28	29	Encephalopathy degree*	0.000000
29	30	Direct Bilirubin (mg/dL)	0.000000
30	31	Diabetes	0.000000
31	32	Creatinine (mg/dL)	0.000000
32	33	Chronic Renal Insufficiency	0.000000
33	34	Arterial Hypertension	0.000000
34	35	Alcohol	0.000000
35	36	Symptoms	0.000000

XGB

XGB0 In CV0...

Checking if correct model is loaded...

XGBClassifier(alpha=0.0002575842389979265, base_score=0.5, booster='gbtree', callbacks=None, colsample_bylevel=1, colsample_bynode=1, colsample_bytree=0.9181376162919086, early_stopping_rounds=None, enable_categorical=False, eta=5.623331491160975e-07, eval_metric=None, gamma=0.0002786718840103683, gpu_id=-1, grow_policy='lossguide', importance_type=None, interaction_constraints='', learning_rate=5.62333128e-07, max_bin=256, max_cat_to_onehot=4, max_delta_step=0, max_depth=27, max_leaves=0, min_child_weight=0.20525460238584922, min_samples_leaf=27, min_samples_split=37, missing=nan, monotone_constraints=('', n_estimators=164, n_jobs=1, nthread=1, ...)

['Alanine transaminase (U/L)', 'Albumin (mg/dL)', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/mL)', 'Arterial Hypertension', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Chronic Renal Insufficiency', 'Cirrhosis', 'Creatinine (mg/dL)', 'Diabetes', 'Direct Bilirubin (mg/dL)', 'Encephalopathy degree*', 'Endemic Countries', 'Esophageal Varices', 'Ferritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haemoglobin (g/dL)', 'Hemochromatosis', 'Hepatitis B Surface Antigen', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibody', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Number of Nodules', 'Obesity', 'Oxygen Saturation (%)', 'Packs of cigarettes per year', 'Performance Status*', 'Portal Hypertension', 'Portal Vein Thrombosis', 'Smoking', 'Splenomegaly', 'Symptoms ', 'Total Bilirubin(mg/dL)']

['Alanine transaminase (U/L)', 'Albumin (mg/dL)', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/mL)', 'Arterial Hypertension', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Chronic Renal Insufficiency', 'Cirrhosis', 'Creatinine (mg/dL)', 'Diabetes', 'Direct Bilirubin (mg/dL)', 'Encephalopathy degree*', 'Endemic Countries', 'Esophageal Varices', 'Ferritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haemoglobin (g/dL)', 'Hemochromatosis', 'Hepatitis B Surface Antigen', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibody', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Number of Nodules', 'Obesity', 'Oxygen Saturation (%)', 'Packs of cigarettes per year', 'Performance Status*', 'Portal Hypertension', 'Portal Vein Thrombosis', 'Smoking', 'Splenomegaly', 'Symptoms ', 'Total Bilirubin(mg/dL)']

Checking explainer for XGB0...

<shap.explainers._tree.Tree object at 0x7f8916d7b970>

Checking shap values for XGB0...

[[[-3.0952360e-06 -3.7619997e-05 -2.3696571e-06 ... -1.2713954e-07
1.3318062e-07 -2.2155659e-06]
[-1.1003718e-05 1.4350392e-05 2.1088497e-06 ... 6.4105130e-08
4.7997673e-07 2.5433717e-06]
[-3.2040905e-06 1.4657915e-05 1.6045622e-05 ... 2.1734811e-08
1.6514244e-07 -8.6456166e-06]
...
[1.4764141e-06 -2.2817851e-05 -1.9707142e-05 ... 4.7425072e-08
1.5751922e-07 6.8362704e-07]
[1.7882336e-05 9.4924808e-06 9.1785603e-07 ... 2.1734811e-08
-1.7996480e-06 7.2638748e-09]
[-3.0676049e-06 1.4631669e-05 5.5639280e-06 ... 2.1734811e-08
-1.6272652e-06 2.0639069e-07]]

Checking shap plots for XGB0...

Expected value for XGB: 1.0591810450932826e-06

Summary Plot for SHAP Values in Test Set:

SHAP Bar Plot for SHAP Values Test Set:

SHAP Decision Plot for SHAP Values in Test Set:

SHAP Decision Plot for Single-Prediction in Test Set:

Checking feature importance for XGB0...

Checking for list of csv files...

['/Users/jessicakim/Desktop/STREAMLINE/DemoData/Output/hcc_demo/hcc-data_example/model_evaluation/shap_values/testResults/XGB_0_shapFIValues_Test.csv']

Unnamed: 0 Features Importance

0 0 Alpha-Fetoprotein (ng/mL) 5.526044e-05

1 1 Albumin (mg/dL) 2.299603e-05

2 2 Aspartate transaminase (U/L) 1.431348e-05

3 3 Alkaline phosphatase (U/L) 1.356586e-05

4 4 Iron 1.137076e-05

5 5 Ferritin (ng/mL) 7.567609e-06

6 6 Haemoglobin (g/dL) 6.252932e-06

7 7 Alanine transaminase (U/L) 5.070476e-06

8 8 Creatinine (mg/dL) 4.338540e-06

9 9 Liver Metastasis 4.279887e-06

10 10 Major dimension of nodule (cm) 4.061806e-06

11 11 Direct Bilirubin (mg/dL) 3.715439e-06

12 12 Performance Status* 3.382434e-06

13 13 Gamma glutamyl transferase (U/L) 3.182782e-06

14 14 Hepatitis B Surface Antigen 2.496254e-06

15 15 Leukocytes(G/L) 2.010043e-06

16 16 Ascites degree* 1.628681e-06

17 17 Endemic Countries 1.511996e-06

18 18 Oxygen Saturation (%) 1.404626e-06

19 19 Cirrhosis 1.295253e-06

20 20 Esophageal Varices 1.257869e-06

21 21 Packs of cigarettes per year 1.130781e-06

22	22	Total Bilirubin(mg/dL)	1.125176e-06
23	23	International Normalised Ratio*	9.767538e-07
24	24	Number of Nodules	9.484509e-07
25	25	Encephalopathy degree*	8.844902e-07
26	26	Mean Corpuscular Volume	7.139903e-07
27	27	Portal Hypertension	6.614388e-07
28	28	Diabetes	5.943173e-07
29	29	Chronic Renal Insufficiency	5.863626e-07
30	30	Symptoms	5.827764e-07
31	31	Portal Vein Thrombosis	4.872277e-07
32	32	Hepatitis C Virus Antibody	4.559925e-07
33	33	Arterial Hypertension	3.621224e-07
34	34	Splenomegaly	8.086869e-08
35	35	Hemochromatosis	6.765081e-08
36	36	Obesity	0.000000e+00
37	37	Hepatitis B e Antigen	0.000000e+00
38	38	Smoking	0.000000e+00

Out[18]:

Unnamed: 0		Features	Importance
0	0	Alpha-Fetoprotein (ng/mL)	5.526044e-05
1	1	Albumin (mg/dL)	2.299603e-05
2	2	Aspartate transaminase (U/L)	1.431348e-05
3	3	Alkaline phosphatase (U/L)	1.356586e-05
4	4	Iron	1.137076e-05
5	5	Ferritin (ng/mL)	7.567609e-06
6	6	Haemoglobin (g/dL)	6.252932e-06
7	7	Alanine transaminase (U/L)	5.070476e-06
8	8	Creatinine (mg/dL)	4.338540e-06
9	9	Liver Metastasis	4.279887e-06
10	10	Major dimension of nodule (cm)	4.061806e-06
11	11	Direct Bilirubin (mg/dL)	3.715439e-06
12	12	Performance Status*	3.382434e-06
13	13	Gamma glutamyl transferase (U/L)	3.182782e-06
14	14	Hepatitis B Surface Antigen	2.496254e-06
15	15	Leukocytes(G/L)	2.010043e-06
16	16	Ascites degree*	1.628681e-06
17	17	Endemic Countries	1.511996e-06
18	18	Oxygen Saturation (%)	1.404626e-06
19	19	Cirrhosis	1.295253e-06
20	20	Esophageal Varices	1.257869e-06
21	21	Packs of cigarets per year	1.130781e-06
22	22	Total Bilirubin(mg/dL)	1.125176e-06
23	23	International Normalised Ratio*	9.767538e-07
24	24	Number of Nodules	9.484509e-07
25	25	Encephalopathy degree*	8.844902e-07
26	26	Mean Corpuscular Volume	7.139903e-07
27	27	Portal Hypertension	6.614388e-07
28	28	Diabetes	5.943173e-07
29	29	Chronic Renal Insufficiency	5.863626e-07
30	30	Symptoms	5.827764e-07
31	31	Portal Vein Thrombosis	4.872277e-07
32	32	Hepatitis C Virus Antibody	4.559925e-07
33	33	Arterial Hypertension	3.621224e-07
34	34	Splenomegaly	8.086869e-08
35	35	Hemochromatosis	6.765081e-08
36	36	Obesity	0.000000e+00
37	37	Hepatitis B e Antigen	0.000000e+00
38	38	Smoking	0.000000e+00

XGB1 In CV1...

```
Checking if correct model is loaded...
XGBClassifier(alpha=0.00029260435288728723, base_score=0.5, booster='gbtree',
              callbacks=None, colsample_bylevel=1, colsample_bynode=1,
              colsample_bytree=0.5441411005619007, early_stopping_rounds=None,
              enable_categorical=False, eta=0.05120369776687421,
              eval_metric=None, gamma=0.4526660690706259, gpu_id=-1,
              grow_policy='depthwise', importance_type=None,
              interaction_constraints='', learning_rate=0.0512036979,
              max_bin=256, max_cat_to_onehot=4, max_delta_step=0, max_depth=18,
              max_leaves=0, min_child_weight=0.12415100550271539,
              min_samples_leaf=9, min_samples_split=27, missing=nan,
              monotone_constraints=('', n_estimators=464, n_jobs=1, nthread=1, ...)
['Age at diagnosis', 'Alanine transaminase (U/L)', 'Albumin (mg/dL)', 'Alcohol', 'Alkaline phosphatase (U/L)', 'Alpha-
Fetoprotein (ng/mL)', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Cirrhosis', 'Creatinine (mg/dL)', 'Diabete
s', 'Direct Bilirubin (mg/dL)', 'Endemic Countries', 'Esophageal Varices', 'Ferritin (ng/mL)', 'Gamma glutamyl transfe
rase (U/L)', 'Grams of Alcohol per day', 'Haemoglobin (g/dL)', 'Hepatitis B Core Antibody', 'Hepatitis B Surface Antig
en', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibody', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/
L)', 'Liver Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Nonalcoholic Steatohepatitis',
'Number of Nodules', 'Obesity', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Platele
ts', 'Portal Hypertension', 'Portal Vein Thrombosis', 'Smoking', 'Symptoms ', 'Total Bilirubin(mg/dL)', 'Total Protein
s (g/dL)']
['Age at diagnosis', 'Alanine transaminase (U/L)', 'Albumin (mg/dL)', 'Alcohol', 'Alkaline phosphatase (U/L)', 'Alpha-
Fetoprotein (ng/mL)', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Cirrhosis', 'Creatinine (mg/dL)', 'Diabete
s', 'Direct Bilirubin (mg/dL)', 'Endemic Countries', 'Esophageal Varices', 'Ferritin (ng/mL)', 'Gamma glutamyl transfe
rase (U/L)', 'Grams of Alcohol per day', 'Haemoglobin (g/dL)', 'Hepatitis B Core Antibody', 'Hepatitis B Surface Antig
en', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibody', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/
L)', 'Liver Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Nonalcoholic Steatohepatitis',
'Number of Nodules', 'Obesity', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Platele
ts', 'Portal Hypertension', 'Portal Vein Thrombosis', 'Smoking', 'Symptoms ', 'Total Bilirubin(mg/dL)', 'Total Protein
s (g/dL)']
```

Checking explainer for XGB1...
<shap.explainers._tree.Tree object at 0x7f88b115c5b0>

Checking shap values for XGB1...

```
[[-0.45429307 -0.06374221 -0.94646686 ... 0.2016137 -0.11681356
-0.14071724]
 [ 0.5519699  0.13245122 -0.10298917 ... -0.83697766 -0.3408652
-0.07272914]
 [ 0.21548487 -0.07396804  0.14945313 ... -0.72553927  0.11201834
 0.11979318]
 ...
 [-0.237244 -0.04841679 -0.3906867 ... -0.8205372 -0.11913119
-0.15298116]
 [-0.33601356 -0.06487641  0.14479543 ... 0.26082134  0.02685894
-0.13012125]
 [ 0.47213364 -0.15212956 -0.5969934 ... 0.20478216 -0.17158583
-0.18285887]]
```

Checking shap plots for XGB1...

Expected value for XGB: 1.5605792999267578
Summary Plot for SHAP Values in Test Set:

SHAP Bar Plot for SHAP Values Test Set:

SHAP Decision Plot for SHAP Values in Test Set:

SHAP Decision Plot for Single-Prediction in Test Set:

Checking feature importance for XGB1...

Checking for list of csv files...

```
['/Users/jessicakim/Desktop/STREAMLINE/DemoData/Output/hcc_demo/hcc-data_example/model_evaluation/shap_values/testResu
lts/XGB_1_shapFIValues_Test.csv']
```

Unnamed: 0	Features	Importance
0	Haemoglobin (g/dL)	0.941411
1	Oxygen Saturation (%)	0.764464
2	Alpha-Fetoprotein (ng/mL)	0.583439
3	Alkaline phosphatase (U/L)	0.477733
4	International Normalised Ratio*	0.449077
5	Direct Bilirubin (mg/dL)	0.401113
6	Symptoms	0.397401
7	Mean Corpuscular Volume	0.392735
8	Ferritin (ng/mL)	0.362923
9	Age at diagnosis	0.357617
10	Albumin (mg/dL)	0.353970
11	Performance Status*	0.301052
12	Leukocytes(G/L)	0.292949
13	Iron	0.286994
14	Aspartate transaminase (U/L)	0.225414
15	Total Bilirubin(mg/dL)	0.177182
16	Major dimension of nodule (cm)	0.176267
17	Ascites degree*	0.174041
18	Grams of Alcohol per day	0.142837
19	Total Proteins (g/dL)	0.139165
20	Platelets	0.116178

21	21	Alanine transaminase (U/L)	0.111249
22	22	Packs of cigarets per year	0.109458
23	23	Creatinine (mg/dL)	0.093624
24	24	Alcohol	0.077872
25	25	Gamma glutamyl transferase (U/L)	0.062831
26	26	Endemic Countries	0.048008
27	27	Number of Nodules	0.047332
28	28	Hepatitis C Virus Antibody	0.020112
29	29	Portal Hypertension	0.019672
30	30	Liver Metastasis	0.013425
31	31	Cirrhosis	0.009383
32	32	Hepatitis B Core Antibody	0.008402
33	33	Diabetes	0.004153
34	34	Hepatitis B Surface Antigen	0.000000
35	35	Obesity	0.000000
36	36	Portal Vein Thrombosis	0.000000
37	37	Esophageal Varices	0.000000
38	38	Smoking	0.000000
39	39	Nonalcoholic Steatohepatitis	0.000000
40	40	Hepatitis B e Antigen	0.000000

Out[18]:

Unnamed: 0		Features	Importance
0	0	Haemoglobin (g/dL)	0.941411
1	1	Oxygen Saturation (%)	0.764464
2	2	Alpha-Fetoprotein (ng/mL)	0.583439
3	3	Alkaline phosphatase (U/L)	0.477733
4	4	International Normalised Ratio*	0.449077
5	5	Direct Bilirubin (mg/dL)	0.401113
6	6	Symptoms	0.397401
7	7	Mean Corpuscular Volume	0.392735
8	8	Ferritin (ng/mL)	0.362923
9	9	Age at diagnosis	0.357617
10	10	Albumin (mg/dL)	0.353970
11	11	Performance Status*	0.301052
12	12	Leukocytes(G/L)	0.292949
13	13	Iron	0.286994
14	14	Aspartate transaminase (U/L)	0.225414
15	15	Total Bilirubin(mg/dL)	0.177182
16	16	Major dimension of nodule (cm)	0.176267
17	17	Ascites degree*	0.174041
18	18	Grams of Alcohol per day	0.142837
19	19	Total Proteins (g/dL)	0.139165
20	20	Platelets	0.116178
21	21	Alanine transaminase (U/L)	0.111249
22	22	Packs of cigarets per year	0.109458
23	23	Creatinine (mg/dL)	0.093624
24	24	Alcohol	0.077872
25	25	Gamma glutamyl transferase (U/L)	0.062831
26	26	Endemic Countries	0.048008
27	27	Number of Nodules	0.047332
28	28	Hepatitis C Virus Antibody	0.020112
29	29	Portal Hypertension	0.019672
30	30	Liver Metastasis	0.013425
31	31	Cirrhosis	0.009383
32	32	Hepatitis B Core Antibody	0.008402
33	33	Diabetes	0.004153
34	34	Hepatitis B Surface Antigen	0.000000
35	35	Obesity	0.000000
36	36	Portal Vein Thrombosis	0.000000
37	37	Esophageal Varices	0.000000
38	38	Smoking	0.000000
39	39	Nonalcoholic Steatohepatitis	0.000000
40	40	Hepatitis B e Antigen	0.000000

XGB2 In CV2...

```
Checking if correct model is loaded...
XGBClassifier(alpha=5.77534955247629e-07, base_score=0.5, booster='gbtree',
              callbacks=None, colsample_bylevel=1, colsample_bynode=1,
              colsample_bytree=0.41771820514444086, early_stopping_rounds=None,
              enable_categorical=False, eta=8.67291826605322e-06,
              eval_metric=None, gamma=0.07212410933578818, gpu_id=-1,
              grow_policy='lossguide', importance_type=None,
              interaction_constraints='', learning_rate=8.67291874e-06,
              max_bin=256, max_cat_to_onehot=4, max_delta_step=0, max_depth=22,
              max_leaves=0, min_child_weight=6.66045104839759,
              min_samples_leaf=11, min_samples_split=39, missing=nan,
              monotone_constraints=('', n_estimators=884, n_jobs=1, nthread=1, ...)
['Age at diagnosis', 'Albumin (mg/dL)', 'Alcohol', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/mL)', 'Arteria
l Hypertension', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Chronic Renal Insufficiency', 'Creatinine (mg/d
L)', 'Diabetes', 'Direct Bilirubin (mg/dL)', 'Encephalopathy degree*', 'Endemic Countries', 'Ferritin (ng/mL)', 'Gamma
glutamyl transferase (U/L)', 'Haemoglobin (g/dL)', 'Hepatitis B Core Antibody', 'Hepatitis B e Antigen', 'Hepatitis C
Virus Antibody', 'Human Immunodeficiency Virus', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver
Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Number of Nodules', 'Obesity', 'Oxygen Satu
ration (%)', 'Packs of cigarets per year', 'Performance Status*', 'Platelets', 'Portal Hypertension', 'Portal Vein Thr
ombosis', 'Symptoms ']
```

Checking explainer for XGB2...
<shap.explainers._tree.Tree object at 0x7f8906b6e3a0>

Checking shap values for XGB2...

```
[[-1.8894803e-04  2.3255567e-04  0.0000000e+00 ...  0.0000000e+00
 -8.9294117e-06 -7.9064384e-06]
 [-1.9001741e-04  1.9959988e-04  0.0000000e+00 ...  0.0000000e+00
 -1.0370755e-05  3.5941350e-06]
 [-1.6012945e-04  1.6149672e-04  0.0000000e+00 ...  0.0000000e+00
 -1.5008896e-05  3.5941350e-06]
 ...
 [-1.7405280e-04  2.5253580e-04  0.0000000e+00 ...  0.0000000e+00
 -1.0370755e-05  3.5941350e-06]
 [-1.8333328e-04 -4.4896262e-04  0.0000000e+00 ...  0.0000000e+00
 -1.5008896e-05  3.5941350e-06]
 [-1.6480772e-04  2.5592663e-04  0.0000000e+00 ...  0.0000000e+00
 -1.3567553e-05  3.5941350e-06]]
```

Checking shap plots for XGB2...

Expected value for XGB: -8.377160702366382e-05
Summary Plot for SHAP Values in Test Set:

SHAP Bar Plot for SHAP Values Test Set:

SHAP Decision Plot for SHAP Values in Test Set:

SHAP Decision Plot for Single-Prediction in Test Set:

Checking feature importance for XGB2...

Checking for list of csv files...

```
['/Users/jessicakim/Desktop/STREAMLINE/DemoData/Output/hcc_demo/hcc-data_example/model_evaluation/shap_values/testResu
lts/XGB_2_shapFIValues_Test.csv']
```

Unnamed: 0	Features	Importance
0	Alkaline phosphatase (U/L)	0.002665
1	Alpha-Fetoprotein (ng/mL)	0.000905
2	Ferritin (ng/mL)	0.000649
3	Haemoglobin (g/dL)	0.000573
4	Iron	0.000544
5	Performance Status*	0.000469
6	Gamma glutamyl transferase (U/L)	0.000292
7	Albumin (mg/dL)	0.000283
8	Leukocytes(G/L)	0.000283
9	Major dimension of nodule (cm)	0.000225
10	Age at diagnosis	0.000182
11	Platelets	0.000104
12	Oxygen Saturation (%)	0.000080
13	Aspartate transaminase (U/L)	0.000069
14	Number of Nodules	0.000040
15	Packs of cigarets per year	0.000036
16	Mean Corpuscular Volume	0.000035
17	Creatinine (mg/dL)	0.000026
18	International Normalised Ratio*	0.000021
19	Diabetes	0.000016
20	Direct Bilirubin (mg/dL)	0.000014
21	Portal Vein Thrombosis	0.000014
22	Liver Metastasis	0.000012

23	23	Symptoms	0.000005
24	24	Encephalopathy degree*	0.000000
25	25	Endemic Countries	0.000000
26	26	Hepatitis B e Antigen	0.000000
27	27	Chronic Renal Insufficiency	0.000000
28	28	Hepatitis C Virus Antibody	0.000000
29	29	Ascites degree*	0.000000
30	30	Arterial Hypertension	0.000000
31	31	Human Immunodeficiency Virus	0.000000
32	32	Obesity	0.000000
33	33	Alcohol	0.000000
34	34	Portal Hypertension	0.000000
35	35	Hepatitis B Core Antibody	0.000000

Out [18]:

Unnamed: 0		Features	Importance
0	0	Alkaline phosphatase (U/L)	0.002665
1	1	Alpha-Fetoprotein (ng/mL)	0.000905
2	2	Ferritin (ng/mL)	0.000649
3	3	Haemoglobin (g/dL)	0.000573
4	4	Iron	0.000544
5	5	Performance Status*	0.000469
6	6	Gamma glutamyl transferase (U/L)	0.000292
7	7	Albumin (mg/dL)	0.000283
8	8	Leukocytes(G/L)	0.000283
9	9	Major dimension of nodule (cm)	0.000225
10	10	Age at diagnosis	0.000182
11	11	Platelets	0.000104
12	12	Oxygen Saturation (%)	0.000080
13	13	Aspartate transaminase (U/L)	0.000069
14	14	Number of Nodules	0.000040
15	15	Packs of cigarets per year	0.000036
16	16	Mean Corpuscular Volume	0.000035
17	17	Creatinine (mg/dL)	0.000026
18	18	International Normalised Ratio*	0.000021
19	19	Diabetes	0.000016
20	20	Direct Bilirubin (mg/dL)	0.000014
21	21	Portal Vein Thrombosis	0.000014
22	22	Liver Metastasis	0.000012
23	23	Symptoms	0.000005
24	24	Encephalopathy degree*	0.000000
25	25	Endemic Countries	0.000000
26	26	Hepatitis B e Antigen	0.000000
27	27	Chronic Renal Insufficiency	0.000000
28	28	Hepatitis C Virus Antibody	0.000000
29	29	Ascites degree*	0.000000
30	30	Arterial Hypertension	0.000000
31	31	Human Immunodeficiency Virus	0.000000
32	32	Obesity	0.000000
33	33	Alcohol	0.000000
34	34	Portal Hypertension	0.000000
35	35	Hepatitis B Core Antibody	0.000000

hcc-data_example_no_covariates

```
{'Symptoms ': 0, 'Alcohol': 1, 'Hepatitis B Surface Antigen': 2, 'Hepatitis B e Antigen': 3, 'Hepatitis B Core Antibod
y': 4, 'Hepatitis C Virus Antibody': 5, 'Cirrhosis': 6, 'Endemic Countries': 7, 'Smoking': 8, 'Diabetes': 9, 'Obesit
y': 10, 'Hemochromatosis': 11, 'Arterial Hypertension': 12, 'Chronic Renal Insufficiency': 13, 'Human Immunodeficiency
Virus': 14, 'Nonalcoholic Steatohepatitis': 15, 'Esophageal Varices': 16, 'Splenomegaly': 17, 'Portal Hypertension': 1
8, 'Portal Vein Thrombosis': 19, 'Liver Metastasis': 20, 'Radiological Hallmark': 21, 'Grams of Alcohol per day': 22,
'Packs of cigarets per year': 23, 'Performance Status*': 24, 'Encephalopathy degree*': 25, 'Ascites degree*': 26, 'Int
ernational Normalised Ratio*': 27, 'Alpha-Fetoprotein (ng/mL)': 28, 'Haemoglobin (g/dL)': 29, 'Mean Corpuscular Volum
e': 30, 'Leukocytes(G/L)': 31, 'Platelets': 32, 'Albumin (mg/dL)': 33, 'Total Bilirubin(mg/dL)': 34, 'Alanine transami
nase (U/L)': 35, 'Aspartate transaminase (U/L)': 36, 'Gamma glutamyl transferase (U/L)': 37, 'Alkaline phosphatase (U/
L)': 38, 'Total Proteins (g/dL)': 39, 'Creatinine (mg/dL)': 40, 'Number of Nodules': 41, 'Major dimension of nodule (c
m)': 42, 'Direct Bilirubin (mg/dL)': 43, 'Iron': 44, 'Oxygen Saturation (%)': 45, 'Ferritin (ng/mL)': 46}
NB
NB0 In CV0...
```

Checking if correct model is loaded...

```
GaussianNB()
['Alanine transaminase (U/L)', 'Albumin (mg/dL)', 'Alcohol', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/m
L)', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Chronic Renal Insufficiency', 'Cirrhosis', 'Creatinine (mg/d
L)', 'Diabetes', 'Direct Bilirubin (mg/dL)', 'Encephalopathy degree*', 'Endemic Countries', 'Esophageal Varices', 'Fer
ritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haemoglobin (g/dL)', 'Hepatitis B Core Antibody', 'Hepatitis B Su
rface Antigen', 'Hepatitis C Virus Antibody', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Met
astasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Nonalcoholic Steatohepatitis', 'Number of Nodu
les', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Portal Vein Thrombosis', 'Splenom
egaly', 'Symptoms ', 'Total Bilirubin(mg/dL)', 'Total Proteins (g/dL)']
['Alanine transaminase (U/L)', 'Albumin (mg/dL)', 'Alcohol', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/m
L)', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Chronic Renal Insufficiency', 'Cirrhosis', 'Creatinine (mg/d
L)', 'Diabetes', 'Direct Bilirubin (mg/dL)', 'Encephalopathy degree*', 'Endemic Countries', 'Esophageal Varices', 'Fer
ritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haemoglobin (g/dL)', 'Hepatitis B Core Antibody', 'Hepatitis B Su
rface Antigen', 'Hepatitis C Virus Antibody', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Met
astasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Nonalcoholic Steatohepatitis', 'Number of Nodu
les', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Portal Vein Thrombosis', 'Splenom
egaly', 'Symptoms ', 'Total Bilirubin(mg/dL)', 'Total Proteins (g/dL)']
```

Checking explainer for NB0...

shap.explainers.Permutation()

Checking shap values for NB0...

```
.values =
array([[ 0.0025      , -0.02      , 0.      , ..., 0.005      ,
        -0.02583333, -0.03      ],
       [ 0.0025      , 0.0125     , 0.      , ..., 0.025      ,
        -0.01833333, -0.0225     ],
       [ 0.01083333, 0.02833333, 0.00166667, ..., 0.05166667,
        -0.01583333, -0.02166667],
       ...,
       [ 0.00333333, -0.01166667, 0.      , ..., 0.005      ,
        -0.02666667, -0.03083333],
       [-0.00083333, -0.005      , 0.      , ..., -0.01916667,
        -0.02583333, -0.03      ],
       [ 0.00666667, 0.02833333, -0.00083333, ..., -0.11333333,
        0.46083333, -0.02333333]])

.base_values =
array([0.32, 0.32, 0.32, 0.32, 0.32, 0.32, 0.32, 0.32, 0.32, 0.32, 0.32,
       0.32, 0.32, 0.32, 0.32, 0.32, 0.32, 0.32, 0.32, 0.32, 0.32, 0.32,
       0.32, 0.32, 0.32, 0.32, 0.32, 0.32, 0.32, 0.32, 0.32, 0.32, 0.32,
       0.32, 0.32, 0.32, 0.32, 0.32, 0.32, 0.32, 0.32, 0.32, 0.32, 0.32])

.data =
array([[ 0.0376161,  1.0547612, -1.7113069, ...,  0.6264224, -0.4385377,
        -0.1829094],
       [-0.331668 , -0.2723382, -1.7113069, ...,  0.6264224, -0.3902633,
        -0.3086845],
       [-0.096669 ,  0.3174837, -1.7113069, ...,  0.6264224, -0.486812 ,
        -0.1829094],
       ...,
       [-0.667381 ,  0.6123947,  0.5843487, ...,  0.6264224, -0.3741719,
        -0.1290058],
       [-0.9191656,  0.3174837,  0.5843487, ..., -1.5963668, -0.2615318,
        -0.1379898],
       [ 0.13833  , -0.346066 ,  0.5843487, ..., -1.5963668,  1.154515 ,
        -0.1379898]])
```

Checking shap plots for NB0...

Summary Plot for SHAP Values in Class 0 & 1 in Test Set:

SHAP Beeswarm Plot for Top 5 SHAP Values in Class 0 & 1 in Test Set:

Checking feature importance for NB0...

Checking for list of csv files...

```
['/Users/jessicakim/Desktop/STREAMLINE/DemoData/Output/hcc_demo/hcc-data_example_no_covariates/model_evaluation/shap_v
alues/testResults/NB_0_shapFIValues_Test.csv']
Unnamed: 0                                Features  Importance
```


0	0	Creatinine (mg/dL)	0.073924
1	1	Encephalopathy degree*	0.062030
2	2	Total Proteins (g/dL)	0.054348
3	3	Total Bilirubin(mg/dL)	0.040697
4	4	Direct Bilirubin (mg/dL)	0.039818
5	5	Performance Status*	0.036258
6	6	Portal Vein Thrombosis	0.032303
7	7	Liver Metastasis	0.028409
8	8	Ascites degree*	0.028379
9	9	Aspartate transaminase (U/L)	0.026303
10	10	Chronic Renal Insufficiency	0.024818
11	11	International Normalised Ratio*	0.023439
12	12	Packs of cigarets per year	0.022970
13	13	Symptoms	0.018394
14	14	Albumin (mg/dL)	0.017955
15	15	Iron	0.017848
16	16	Alpha-Fetoprotein (ng/mL)	0.017712
17	17	Major dimension of nodule (cm)	0.016288
18	18	Esophageal Varices	0.016106
19	19	Endemic Countries	0.013924
20	20	Diabetes	0.013758
21	21	Haemoglobin (g/dL)	0.013485
22	22	Oxygen Saturation (%)	0.013288
23	23	Leukocytes(G/L)	0.012621
24	24	Hepatitis C Virus Antibody	0.011561
25	25	Hepatitis B Surface Antigen	0.011121
26	26	Alkaline phosphatase (U/L)	0.010121
27	27	Gamma glutamyl transferase (U/L)	0.009909
28	28	Nonalcoholic Steatohepatitis	0.008152
29	29	Ferritin (ng/mL)	0.007939
30	30	Cirrhosis	0.007167
31	31	Alanine transaminase (U/L)	0.005561
32	32	Hepatitis B Core Antibody	0.005455
33	33	Mean Corpuscular Volume	0.005197
34	34	Number of Nodules	0.003439
35	35	Splenomegaly	0.002000
36	36	Alcohol	0.000121

Out [18]:

Unnamed: 0		Features	Importance
0	0	Creatinine (mg/dL)	0.073924
1	1	Encephalopathy degree*	0.062030
2	2	Total Proteins (g/dL)	0.054348
3	3	Total Bilirubin(mg/dL)	0.040697
4	4	Direct Bilirubin (mg/dL)	0.039818
5	5	Performance Status*	0.036258
6	6	Portal Vein Thrombosis	0.032303
7	7	Liver Metastasis	0.028409
8	8	Ascites degree*	0.028379
9	9	Aspartate transaminase (U/L)	0.026303
10	10	Chronic Renal Insufficiency	0.024818
11	11	International Normalised Ratio*	0.023439
12	12	Packs of cigarets per year	0.022970
13	13	Symptoms	0.018394
14	14	Albumin (mg/dL)	0.017955
15	15	Iron	0.017848
16	16	Alpha-Fetoprotein (ng/mL)	0.017712
17	17	Major dimension of nodule (cm)	0.016288
18	18	Esophageal Varices	0.016106
19	19	Endemic Countries	0.013924
20	20	Diabetes	0.013758
21	21	Haemoglobin (g/dL)	0.013485
22	22	Oxygen Saturation (%)	0.013288
23	23	Leukocytes(G/L)	0.012621
24	24	Hepatitis C Virus Antibody	0.011561
25	25	Hepatitis B Surface Antigen	0.011121
26	26	Alkaline phosphatase (U/L)	0.010121
27	27	Gamma glutamyl transferase (U/L)	0.009909
28	28	Nonalcoholic Steatohepatitis	0.008152
29	29	Ferritin (ng/mL)	0.007939
30	30	Cirrhosis	0.007167
31	31	Alanine transaminase (U/L)	0.005561
32	32	Hepatitis B Core Antibody	0.005455
33	33	Mean Corpuscular Volume	0.005197
34	34	Number of Nodules	0.003439
35	35	Splenomegaly	0.002000
36	36	Alcohol	0.000121

NB1 In CV1...

```
Checking if correct model is loaded...
GaussianNB()
['Alanine transaminase (U/L)', 'Albumin (mg/dL)', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/mL)', 'Arterial Hypertension', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Cirrhosis', 'Creatinine (mg/dL)', 'Diabetes', 'Direct Bilirubin (mg/dL)', 'Encephalopathy degree*', 'Endemic Countries', 'Ferritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haemoglobin (g/dL)', 'Hemochromatosis', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibody', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Nonalcoholic Steatohepatitis', 'Number of Nodules', 'Obesity', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Platelets', 'Portal Hypertension', 'Portal Vein Thrombosis', 'Symptoms ', 'Total Bilirubin(mg/dL)']
['Alanine transaminase (U/L)', 'Albumin (mg/dL)', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/mL)', 'Arterial Hypertension', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Cirrhosis', 'Creatinine (mg/dL)', 'Diabetes', 'Direct Bilirubin (mg/dL)', 'Encephalopathy degree*', 'Endemic Countries', 'Ferritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haemoglobin (g/dL)', 'Hemochromatosis', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibody', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Nonalcoholic Steatohepatitis', 'Number of Nodules', 'Obesity', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Platelets', 'Portal Hypertension', 'Portal Vein Thrombosis', 'Symptoms ', 'Total Bilirubin(mg/dL)']
```

Checking explainer for NB1...
shap.explainers.Permutation()

Checking shap values for NB1...

```
.values =
array([[ 0.0225      , -0.10916667,  0.47666667, ..., -0.02083333,
         0.0825      , -0.005       ],
       [-0.00416667, -0.00916667, -0.05416667, ..., -0.08083333,
        -0.22833333, -0.025       ],
       [ 0.0075      ,  0.025       ,  0.05666667, ..., -0.0025      ,
        -0.06833333, -0.01083333],
       ...,
       [-0.04666667, -0.04416667,  0.145       , ..., -0.01833333,
        -0.25       , -0.03833333],
       [-0.0025      ,  0.00666667,  0.01666667, ...,  0.         ,
         0.01083333,  0.         ],
       [ 0.01083333, -0.02       ,  0.01       , ..., -0.02333333,
         0.0975      , -0.01416667]])

.base_values =
array([0.48, 0.48, 0.48, 0.48, 0.48, 0.48, 0.48, 0.48, 0.48, 0.48, 0.48,
       0.48, 0.48, 0.48, 0.48, 0.48, 0.48, 0.48, 0.48, 0.48, 0.48, 0.48,
       0.48, 0.48, 0.48, 0.48, 0.48, 0.48, 0.48, 0.48, 0.48, 0.48, 0.48,
       0.48, 0.48, 0.48, 0.48, 0.48, 0.48, 0.48, 0.48, 0.48, 0.48])

.data =
array([[ -0.3307946,  1.9018224,  2.7240451, ..., -0.5141671,  0.6831301,
        -0.4175834],
       [-0.9380037,  0.0449382, -0.5189802, ..., -0.5141671, -1.4638501,
        -0.449261 ],
       [-0.2159173, -0.5585492,  0.9323539, ..., -0.5141671, -1.4638501,
        -0.1800009],
       ...,
       [ 0.8343903,  0.1996785,  1.465366 , ..., -0.5141671, -1.4638501,
        -0.3700669],
       [ 1.1462003, -0.1098022,  1.3626167, ...,  1.944893 ,  0.6831301,
        -0.2433562],
       [-0.6754268,  1.1281206,  0.1103594, ..., -0.5141671,  0.6831301,
        -0.3859057]])
```

Checking shap plots for NB1...

Summary Plot for SHAP Values in Class 0 & 1 in Test Set:

SHAP Beeswarm Plot for Top 5 SHAP Values in Class 0 & 1 in Test Set:

Checking feature importance for NB1...

Checking for list of csv files...

```
['/Users/jessicakim/Desktop/STREAMLINE/DemoData/Output/hcc_demo/hcc-data_example_no_covariates/model_evaluation/shap_v
alues/testResults/NB_1_shapFIVValues_Test.csv']
```

Unnamed: 0	Features	Importance
0	Endemic Countries	0.095621
1	Symptoms	0.069697
2	Nonalcoholic Steatohepatitis	0.065742
3	Performance Status*	0.056652
4	Ascites degree*	0.055470
5	Alkaline phosphatase (U/L)	0.049424
6	Albumin (mg/dL)	0.042136
7	International Normalised Ratio*	0.040924
8	Portal Hypertension	0.040121
9	Haemoglobin (g/dL)	0.037000
10	Portal Vein Thrombosis	0.027621
11	Liver Metastasis	0.026121
12	Alpha-Fetoprotein (ng/mL)	0.025894
13	Total Bilirubin(mg/dL)	0.025318
14	Encephalopathy degree*	0.024485

15	15	Direct Bilirubin (mg/dL)	0.023394
16	16	Iron	0.023212
17	17	Oxygen Saturation (%)	0.022545
18	18	Platelets	0.020167
19	19	Ferritin (ng/mL)	0.015409
20	20	Mean Corpuscular Volume	0.014773
21	21	Packs of cigarets per year	0.014652
22	22	Alanine transaminase (U/L)	0.014106
23	23	Major dimension of nodule (cm)	0.012758
24	24	Hepatitis C Virus Antibody	0.010076
25	25	Diabetes	0.008015
26	26	Aspartate transaminase (U/L)	0.007818
27	27	Obesity	0.006606
28	28	Arterial Hypertension	0.005636
29	29	Hemochromatosis	0.005500
30	30	Gamma glutamyl transferase (U/L)	0.005455
31	31	Creatinine (mg/dL)	0.003591
32	32	Number of Nodules	0.003045
33	33	Hepatitis B e Antigen	0.002303
34	34	Leukocytes(G/L)	0.000318
35	35	Cirrhosis	0.000000

Out [18]:

Unnamed: 0		Features	Importance
0	0	Endemic Countries	0.095621
1	1	Symptoms	0.069697
2	2	Nonalcoholic Steatohepatitis	0.065742
3	3	Performance Status*	0.056652
4	4	Ascites degree*	0.055470
5	5	Alkaline phosphatase (U/L)	0.049424
6	6	Albumin (mg/dL)	0.042136
7	7	International Normalised Ratio*	0.040924
8	8	Portal Hypertension	0.040121
9	9	Haemoglobin (g/dL)	0.037000
10	10	Portal Vein Thrombosis	0.027621
11	11	Liver Metastasis	0.026121
12	12	Alpha-Fetoprotein (ng/mL)	0.025894
13	13	Total Bilirubin(mg/dL)	0.025318
14	14	Encephalopathy degree*	0.024485
15	15	Direct Bilirubin (mg/dL)	0.023394
16	16	Iron	0.023212
17	17	Oxygen Saturation (%)	0.022545
18	18	Platelets	0.020167
19	19	Ferritin (ng/mL)	0.015409
20	20	Mean Corpuscular Volume	0.014773
21	21	Packs of cigarets per year	0.014652
22	22	Alanine transaminase (U/L)	0.014106
23	23	Major dimension of nodule (cm)	0.012758
24	24	Hepatitis C Virus Antibody	0.010076
25	25	Diabetes	0.008015
26	26	Aspartate transaminase (U/L)	0.007818
27	27	Obesity	0.006606
28	28	Arterial Hypertension	0.005636
29	29	Hemochromatosis	0.005500
30	30	Gamma glutamyl transferase (U/L)	0.005455
31	31	Creatinine (mg/dL)	0.003591
32	32	Number of Nodules	0.003045
33	33	Hepatitis B e Antigen	0.002303
34	34	Leukocytes(G/L)	0.000318
35	35	Cirrhosis	0.000000

NB2 In CV2...

```
Checking if correct model is loaded...
GaussianNB()
['Albumin (mg/dL)', 'Alcohol', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/mL)', 'Arterial Hypertension', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Chronic Renal Insufficiency', 'Cirrhosis', 'Creatinine (mg/dL)', 'Diabetes', 'Direct Bilirubin (mg/dL)', 'Endemic Countries', 'Ferritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haemoglobin (g/dL)', 'Hemochromatosis', 'Hepatitis B Surface Antigen', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibody', 'Human Immunodeficiency Virus', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Nonalcoholic Steatohepatitis', 'Number of Nodules', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Platelets', 'Portal Hypertension', 'Portal Vein Thrombosis', 'Smoking']
['Albumin (mg/dL)', 'Alcohol', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/mL)', 'Arterial Hypertension', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Chronic Renal Insufficiency', 'Cirrhosis', 'Creatinine (mg/dL)', 'Diabetes', 'Direct Bilirubin (mg/dL)', 'Endemic Countries', 'Ferritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haemoglobin (g/dL)', 'Hemochromatosis', 'Hepatitis B Surface Antigen', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibody', 'Human Immunodeficiency Virus', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Nonalcoholic Steatohepatitis', 'Number of Nodules', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Platelets', 'Portal Hypertension', 'Portal Vein Thrombosis', 'Smoking']
```

Checking explainer for NB2...
shap.explainers.Permutation()

Checking shap values for NB2...

```
.values =
array([[ 0.          ,  0.          ,  0.          , ...,  0.          ,
         0.          ,  0.          ],
       [-0.00083333,  0.          ,  0.0175         , ...,  0.          ,
         0.          ,  0.          ],
       [ 0.00916667, -0.00166667,  0.01083333, ..., -0.00833333,
        -0.00583333,  0.01          ],
       ...,
       [ 0.01833333,  0.          ,  0.0175         , ...,  0.          ,
        -0.00166667, -0.00416667],
       [-0.03083333,  0.005         ,  0.02166667, ..., -0.00166667,
        -0.03416667, -0.02          ],
       [ 0.          ,  0.          ,  0.          , ...,  0.          ,
         0.          ,  0.          ]])

.base_values =
array([0.05, 0.05, 0.05, 0.05, 0.05, 0.05, 0.05, 0.05, 0.05, 0.05, 0.05,
       0.05, 0.05, 0.05, 0.05, 0.05, 0.05, 0.05, 0.05, 0.05, 0.05, 0.05,
       0.05, 0.05, 0.05, 0.05, 0.05, 0.05, 0.05, 0.05, 0.05, 0.05, 0.05,
       0.05, 0.05, 0.05, 0.05, 0.05, 0.05, 0.05, 0.05, 0.05, 0.05])

.data =
array([[ -0.4184767,  0.5703518, -0.3573003, ...,  0.6264224, -0.5282705,
        -0.7859052],
       [ -0.4184767,  0.5703518,  4.7719674, ..., -1.5963668, -0.5282705,
         1.272418 ],
       [ 0.0205491, -1.7533038,  0.3559594, ...,  0.6264224, -0.5282705,
        -0.7859052],
       ...,
       [-1.8818959,  0.5703518,  1.5964111, ...,  0.6264224, -0.5282705,
         1.272418 ],
       [ 1.0449426,  0.5703518, -0.1898393, ...,  0.6264224, -0.5282705,
         1.272418 ],
       [-2.0282378,  0.5703518, -0.5867838, ...,  0.6264224, -0.5282705,
        -0.7859052]])
```

Checking shap plots for NB2...

Summary Plot for SHAP Values in Class 0 & 1 in Test Set:

SHAP Beeswarm Plot for Top 5 SHAP Values in Class 0 & 1 in Test Set:

Checking feature importance for NB2...

Checking for list of csv files...

```
['/Users/jessicakim/Desktop/STREAMLINE/DemoData/Output/hcc_demo/hcc-data_example_no_covariates/model_evaluation/shap_v
alues/testResults/NB_2_shapFIValues_Test.csv']
```

Unnamed: 0	Features	Importance
0	0 Direct Bilirubin (mg/dL)	0.059470
1	1 Human Immunodeficiency Virus	0.053273
2	2 Packs of cigarets per year	0.017803
3	3 International Normalised Ratio*	0.013758
4	4 Ferritin (ng/mL)	0.013424
5	5 Hepatitis B e Antigen	0.009879
6	6 Nonalcoholic Steatohepatitis	0.003909
7	7 Liver Metastasis	0.003712
8	8 Ascites degree*	0.003106
9	9 Aspartate transaminase (U/L)	0.002409
10	10 Leukocytes(G/L)	0.002227
11	11 Platelets	0.001864
12	12 Alkaline phosphatase (U/L)	0.001652
13	13 Hepatitis B Surface Antigen	0.001621
14	14 Performance Status*	0.001576

15	15	Hemochromatosis	0.001576
16	16	Haemoglobin (g/dL)	0.001455
17	17	Albumin (mg/dL)	0.001364
18	18	Portal Vein Thrombosis	0.001136
19	19	Major dimension of nodule (cm)	0.001091
20	20	Mean Corpuscular Volume	0.001076
21	21	Number of Nodules	0.001015
22	22	Hepatitis C Virus Antibody	0.000985
23	23	Gamma glutamyl transferase (U/L)	0.000818
24	24	Diabetes	0.000803
25	25	Iron	0.000803
26	26	Smoking	0.000712
27	27	Arterial Hypertension	0.000652
28	28	Chronic Renal Insufficiency	0.000485
29	29	Cirrhosis	0.000424
30	30	Alpha-Fetoprotein (ng/mL)	0.000409
31	31	Creatinine (mg/dL)	0.000273
32	32	Oxygen Saturation (%)	0.000212
33	33	Portal Hypertension	0.000182
34	34	Alcohol	0.000136
35	35	Endemic Countries	0.000045

Out [18]:

Unnamed: 0		Features	Importance
0	0	Direct Bilirubin (mg/dL)	0.059470
1	1	Human Immunodeficiency Virus	0.053273
2	2	Packs of cigarets per year	0.017803
3	3	International Normalised Ratio*	0.013758
4	4	Ferritin (ng/mL)	0.013424
5	5	Hepatitis B e Antigen	0.009879
6	6	Nonalcoholic Steatohepatitis	0.003909
7	7	Liver Metastasis	0.003712
8	8	Ascites degree*	0.003106
9	9	Aspartate transaminase (U/L)	0.002409
10	10	Leukocytes(G/L)	0.002227
11	11	Platelets	0.001864
12	12	Alkaline phosphatase (U/L)	0.001652
13	13	Hepatitis B Surface Antigen	0.001621
14	14	Performance Status*	0.001576
15	15	Hemochromatosis	0.001576
16	16	Haemoglobin (g/dL)	0.001455
17	17	Albumin (mg/dL)	0.001364
18	18	Portal Vein Thrombosis	0.001136
19	19	Major dimension of nodule (cm)	0.001091
20	20	Mean Corpuscular Volume	0.001076
21	21	Number of Nodules	0.001015
22	22	Hepatitis C Virus Antibody	0.000985
23	23	Gamma glutamyl transferase (U/L)	0.000818
24	24	Diabetes	0.000803
25	25	Iron	0.000803
26	26	Smoking	0.000712
27	27	Arterial Hypertension	0.000652
28	28	Chronic Renal Insufficiency	0.000485
29	29	Cirrhosis	0.000424
30	30	Alpha-Fetoprotein (ng/mL)	0.000409
31	31	Creatinine (mg/dL)	0.000273
32	32	Oxygen Saturation (%)	0.000212
33	33	Portal Hypertension	0.000182
34	34	Alcohol	0.000136
35	35	Endemic Countries	0.000045

```
LR
LR0 In CV0...

Checking if correct model is loaded...
LogisticRegression(C=0.0076324520136090606, dual=True,
                    max_iter=383.683139958808, random_state=42,
                    solver='liblinear')
['Alanine transaminase (U/L)', 'Albumin (mg/dL)', 'Alcohol', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/m
L)', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Chronic Renal Insufficiency', 'Cirrhosis', 'Creatinine (mg/d
L)', 'Diabetes', 'Direct Bilirubin (mg/dL)', 'Encephalopathy degree*', 'Endemic Countries', 'Esophageal Varices', 'Fer
ritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haemoglobin (g/dL)', 'Hepatitis B Core Antibody', 'Hepatitis B Su
rface Antigen', 'Hepatitis C Virus Antibody', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Met
astasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Nonalcoholic Steatohepatitis', 'Number of Nodu
les', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Portal Vein Thrombosis', 'Splenom
egaly', 'Symptoms ', 'Total Bilirubin(mg/dL)', 'Total Proteins (g/dL)']
['Alanine transaminase (U/L)', 'Albumin (mg/dL)', 'Alcohol', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/m
L)', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Chronic Renal Insufficiency', 'Cirrhosis', 'Creatinine (mg/d
L)', 'Diabetes', 'Direct Bilirubin (mg/dL)', 'Encephalopathy degree*', 'Endemic Countries', 'Esophageal Varices', 'Fer
ritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haemoglobin (g/dL)', 'Hepatitis B Core Antibody', 'Hepatitis B Su
rface Antigen', 'Hepatitis C Virus Antibody', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Met
astasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Nonalcoholic Steatohepatitis', 'Number of Nodu
les', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Portal Vein Thrombosis', 'Splenom
egaly', 'Symptoms ', 'Total Bilirubin(mg/dL)', 'Total Proteins (g/dL)']

Checking explainer for LR0...
<shap.explainers._linear.Linear object at 0x7f89160e3340>

Checking shap values for LR0...

[[-0.00015674 -0.05535253  0.01520368 ...  0.02453187 -0.02176526
  -0.01030959]
 [ 0.00068073  0.02080485  0.01520368 ...  0.02453187 -0.01946545
  -0.01559663]
 [ 0.0001478  -0.01304287  0.01520368 ...  0.02453187 -0.02406506
  -0.01030959]
 ...
 [ 0.00144207 -0.02996674 -0.00683064 ...  0.02453187 -0.01869885
  -0.00804372]
 [ 0.00201307 -0.01304287 -0.00683064 ... -0.0663269  -0.01333263
  -0.00842137]
 [-0.00038514  0.02503582 -0.00683064 ... -0.0663269   0.05412839
  -0.00842137]]

Checking shap plots for LR0...

Expected value for LR: -0.032810633342135964
Summary Plot for SHAP Values in Test Set:

SHAP Bar Plot for SHAP Values Test Set:

SHAP Decision Plot for SHAP Values in Test Set:

SHAP Decision Plot for Single-Prediction in Test Set:

Checking feature importance for LR0...

Checking for list of csv files...

['/Users/jessicakim/Desktop/STREAMLINE/DemoData/Output/hcc_demo/hcc-data_example_no_covariates/model_evaluation/shap_v
alues/testResults/LR_0_shapFIValues_Test.csv']
  Unnamed: 0      Features  Importance
0           0      Performance Status*    0.073896
1           1      Liver Metastasis    0.066073
2           2      Portal Vein Thrombosis    0.056833
3           3      Diabetes    0.049721
4           4  Major dimension of nodule (cm)    0.048746
5           5      Aspartate transaminase (U/L)    0.047610
6           6      Ascites degree*    0.046914
7           7      Esophageal Varices    0.046794
8           8      Albumin (mg/dL)    0.045286
9           9      Alkaline phosphatase (U/L)    0.042692
10          10      Symptoms    0.041250
11          11      Ferritin (ng/mL)    0.040764
12          12      Creatinine (mg/dL)    0.039875
13          13      Haemoglobin (g/dL)    0.035179
14          14  Gamma glutamyl transferase (U/L)    0.030540
15          15      Hepatitis C Virus Antibody    0.028889
16          16      Chronic Renal Insufficiency    0.024176
17          17      Leukocytes(G/L)    0.023098
18          18      Hepatitis B Surface Antigen    0.021932
19          19      Encephalopathy degree*    0.019545
20          20      Direct Bilirubin (mg/dL)    0.019318
21          21      Endemic Countries    0.018639
22          22      International Normalised Ratio*    0.018475
23          23      Total Bilirubin(mg/dL)    0.018277
24          24      Total Proteins (g/dL)    0.017360
25          25      Packs of cigarets per year    0.015368
26          26      Hepatitis B Core Antibody    0.011204
27          27      Cirrhosis    0.010850
28          28      Alcohol    0.009114
29          29      Alpha-Fetoprotein (ng/mL)    0.006761
```

30	30	Nonalcoholic Steatohepatitis	0.005990
31	31	Mean Corpuscular Volume	0.003729
32	32	Oxygen Saturation (%)	0.002960
33	33	Iron	0.002790
34	34	Splenomegaly	0.001654
35	35	Alanine transaminase (U/L)	0.001401
36	36	Number of Nodules	0.001181

Out [18]:

Unnamed: 0		Features	Importance
0	0	Performance Status*	0.073896
1	1	Liver Metastasis	0.066073
2	2	Portal Vein Thrombosis	0.056833
3	3	Diabetes	0.049721
4	4	Major dimension of nodule (cm)	0.048746
5	5	Aspartate transaminase (U/L)	0.047610
6	6	Ascites degree*	0.046914
7	7	Esophageal Varices	0.046794
8	8	Albumin (mg/dL)	0.045286
9	9	Alkaline phosphatase (U/L)	0.042692
10	10	Symptoms	0.041250
11	11	Ferritin (ng/mL)	0.040764
12	12	Creatinine (mg/dL)	0.039875
13	13	Haemoglobin (g/dL)	0.035179
14	14	Gamma glutamyl transferase (U/L)	0.030540
15	15	Hepatitis C Virus Antibody	0.028889
16	16	Chronic Renal Insufficiency	0.024176
17	17	Leukocytes(G/L)	0.023098
18	18	Hepatitis B Surface Antigen	0.021932
19	19	Encephalopathy degree*	0.019545
20	20	Direct Bilirubin (mg/dL)	0.019318
21	21	Endemic Countries	0.018639
22	22	International Normalised Ratio*	0.018475
23	23	Total Bilirubin(mg/dL)	0.018277
24	24	Total Proteins (g/dL)	0.017360
25	25	Packs of cigarets per year	0.015368
26	26	Hepatitis B Core Antibody	0.011204
27	27	Cirrhosis	0.010850
28	28	Alcohol	0.009114
29	29	Alpha-Fetoprotein (ng/mL)	0.006761
30	30	Nonalcoholic Steatohepatitis	0.005990
31	31	Mean Corpuscular Volume	0.003729
32	32	Oxygen Saturation (%)	0.002960
33	33	Iron	0.002790
34	34	Splenomegaly	0.001654
35	35	Alanine transaminase (U/L)	0.001401
36	36	Number of Nodules	0.001181

LR1 In CV1...

```
Checking if correct model is loaded...
LogisticRegression(C=0.0003630322466779861, class_weight='balanced',
                    max_iter=159.30522616241012, random_state=42)
['Alanine transaminase (U/L)', 'Albumin (mg/dL)', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/mL)', 'Arterial Hypertension', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Cirrhosis', 'Creatinine (mg/dL)', 'Diabetes', 'Direct Bilirubin (mg/dL)', 'Encephalopathy degree*', 'Endemic Countries', 'Ferritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haemoglobin (g/dL)', 'Hemochromatosis', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibody', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Nonalcoholic Steatohepatitis', 'Number of Nodules', 'Obesity', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Platelets', 'Portal Hypertension', 'Portal Vein Thrombosis', 'Symptoms ', 'Total Bilirubin(mg/dL)']
['Alanine transaminase (U/L)', 'Albumin (mg/dL)', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/mL)', 'Arterial Hypertension', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Cirrhosis', 'Creatinine (mg/dL)', 'Diabetes', 'Direct Bilirubin (mg/dL)', 'Encephalopathy degree*', 'Endemic Countries', 'Ferritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haemoglobin (g/dL)', 'Hemochromatosis', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibody', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Nonalcoholic Steatohepatitis', 'Number of Nodules', 'Obesity', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Platelets', 'Portal Hypertension', 'Portal Vein Thrombosis', 'Symptoms ', 'Total Bilirubin(mg/dL)']
```

Checking explainer for LR1...
<shap.explainers._linear.Linear object at 0x7f891600ad90>

Checking shap values for LR1...

```
[[ 0.00034788 -0.01330814  0.01706337 ... -0.00222993  0.00364155
  -0.00301374]
 [ 0.00099548 -0.00012806 -0.00335686 ... -0.00222993 -0.00984567
  -0.00321245]
 [ 0.00022536  0.00415547  0.0057817  ... -0.00222993 -0.00984567
  -0.00152343]
 ...
 [-0.00089483 -0.0012264   0.00913789 ... -0.00222993 -0.00984567
  -0.00271568]
 [-0.00122738  0.00097028  0.00849092 ...  0.00602908  0.00364155
  -0.00192084]
 [ 0.00071544 -0.00781644  0.00060588 ... -0.00222993  0.00364155
  -0.00281503]]
```

Checking shap plots for LR1...

Expected value for LR: -0.004693505131985552
Summary Plot for SHAP Values in Test Set:

SHAP Bar Plot for SHAP Values Test Set:

SHAP Decision Plot for SHAP Values in Test Set:

SHAP Decision Plot for Single-Prediction in Test Set:

Checking feature importance for LR1...

Checking for list of csv files...

```
['/Users/jessicakim/Desktop/STREAMLINE/DemoData/Output/hcc_demo/hcc-data_example_no_covariates/model_evaluation/shap_v
alues/testResults/LR_1_shapFIValues_Test.csv']
```

Unnamed: 0	0	Features	Importance
0	0	Performance Status*	0.007802
1	1	Ascites degree*	0.007155
2	2	Haemoglobin (g/dL)	0.006427
3	3	Albumin (mg/dL)	0.006404
4	4	International Normalised Ratio*	0.005786
5	5	Symptoms	0.005672
6	6	Alkaline phosphatase (U/L)	0.005199
7	7	Alpha-Fetoprotein (ng/mL)	0.004664
8	8	Iron	0.003656
9	9	Portal Hypertension	0.003202
10	10	Portal Vein Thrombosis	0.003128
11	11	Encephalopathy degree*	0.002886
12	12	Ferritin (ng/mL)	0.002743
13	13	Liver Metastasis	0.002591
14	14	Total Bilirubin(mg/dL)	0.002528
15	15	Major dimension of nodule (cm)	0.002486
16	16	Endemic Countries	0.002159
17	17	Direct Bilirubin (mg/dL)	0.002138
18	18	Nonalcoholic Steatohepatitis	0.002059
19	19	Platelets	0.001929
20	20	Aspartate transaminase (U/L)	0.001460
21	21	Diabetes	0.001386
22	22	Oxygen Saturation (%)	0.001267
23	23	Hepatitis C Virus Antibody	0.001227
24	24	Arterial Hypertension	0.001105
25	25	Mean Corpuscular Volume	0.001055
26	26	Gamma glutamyl transferase (U/L)	0.001006
27	27	Number of Nodules	0.000676
28	28	Alanine transaminase (U/L)	0.000660
29	29	Hepatitis B e Antigen	0.000520
30	30	Creatinine (mg/dL)	0.000488
31	31	Obesity	0.000443

32	32	Packs of cigarets per year	0.000401
33	33	Leukocytes(G/L)	0.000167
34	34	Hemochromatosis	0.000151
35	35	Cirrhosis	0.000024

Out [18]:

Unnamed: 0		Features	Importance
0	0	Performance Status*	0.007802
1	1	Ascites degree*	0.007155
2	2	Haemoglobin (g/dL)	0.006427
3	3	Albumin (mg/dL)	0.006404
4	4	International Normalised Ratio*	0.005786
5	5	Symptoms	0.005672
6	6	Alkaline phosphatase (U/L)	0.005199
7	7	Alpha-Fetoprotein (ng/mL)	0.004664
8	8	Iron	0.003656
9	9	Portal Hypertension	0.003202
10	10	Portal Vein Thrombosis	0.003128
11	11	Encephalopathy degree*	0.002886
12	12	Ferritin (ng/mL)	0.002743
13	13	Liver Metastasis	0.002591
14	14	Total Bilirubin(mg/dL)	0.002528
15	15	Major dimension of nodule (cm)	0.002486
16	16	Endemic Countries	0.002159
17	17	Direct Bilirubin (mg/dL)	0.002138
18	18	Nonalcoholic Steatohepatitis	0.002059
19	19	Platelets	0.001929
20	20	Aspartate transaminase (U/L)	0.001460
21	21	Diabetes	0.001386
22	22	Oxygen Saturation (%)	0.001267
23	23	Hepatitis C Virus Antibody	0.001227
24	24	Arterial Hypertension	0.001105
25	25	Mean Corpuscular Volume	0.001055
26	26	Gamma glutamyl transferase (U/L)	0.001006
27	27	Number of Nodules	0.000676
28	28	Alanine transaminase (U/L)	0.000660
29	29	Hepatitis B e Antigen	0.000520
30	30	Creatinine (mg/dL)	0.000488
31	31	Obesity	0.000443
32	32	Packs of cigarets per year	0.000401
33	33	Leukocytes(G/L)	0.000167
34	34	Hemochromatosis	0.000151
35	35	Cirrhosis	0.000024

LR2 In CV2...

```
Checking if correct model is loaded...
LogisticRegression(C=1.1355889848345609e-05, dual=True,
                    max_iter=287.08753481954665, random_state=42,
                    solver='liblinear')
['Albumin (mg/dL)', 'Alcohol', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/mL)', 'Arterial Hypertension', 'As
cites degree*', 'Aspartate transaminase (U/L)', 'Chronic Renal Insufficiency', 'Cirrhosis', 'Creatinine (mg/dL)', 'Dia
betes', 'Direct Bilirubin (mg/dL)', 'Endemic Countries', 'Ferritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haem
oglobin (g/dL)', 'Hemochromatosis', 'Hepatitis B Surface Antigen', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibod
y', 'Human Immunodeficiency Virus', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Metastasis',
'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Nonalcoholic Steatohepatitis', 'Number of Nodules', 'Oxy
gen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Platelets', 'Portal Hypertension', 'Portal
Vein Thrombosis', 'Smoking']
['Albumin (mg/dL)', 'Alcohol', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/mL)', 'Arterial Hypertension', 'As
cites degree*', 'Aspartate transaminase (U/L)', 'Chronic Renal Insufficiency', 'Cirrhosis', 'Creatinine (mg/dL)', 'Dia
betes', 'Direct Bilirubin (mg/dL)', 'Endemic Countries', 'Ferritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haem
oglobin (g/dL)', 'Hemochromatosis', 'Hepatitis B Surface Antigen', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibod
y', 'Human Immunodeficiency Virus', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Metastasis',
'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Nonalcoholic Steatohepatitis', 'Number of Nodules', 'Oxy
gen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Platelets', 'Portal Hypertension', 'Portal
Vein Thrombosis', 'Smoking']
```

Checking explainer for LR2...
<shap.explainers._linear.Linear object at 0x7f8925b355e0>

Checking shap values for LR2...

```
[[ 7.32083460e-05  1.92822706e-05 -4.36797008e-05 ... -3.15759647e-05
 -7.30478160e-05  3.36532768e-05]
 [ 7.32083460e-05  1.92822706e-05  8.74173101e-04 ...  8.11953378e-05
 -7.30478160e-05 -5.73015253e-05]
 [ 7.45330978e-06 -7.71290826e-05  8.39539991e-05 ... -3.15759647e-05
 -7.30478160e-05  3.36532768e-05]
 ...
 [ 2.92391780e-04  1.92822706e-05  3.05925660e-04 ... -3.15759647e-05
 -7.30478160e-05 -5.73015253e-05]
 [-1.45975103e-04  1.92822706e-05 -1.37135230e-05 ... -3.15759647e-05
 -7.30478160e-05 -5.73015253e-05]
 [ 3.14310121e-04  1.92822706e-05 -8.47444465e-05 ... -3.15759647e-05
 -7.30478160e-05  3.36532768e-05]]
```

Checking shap plots for LR2...

Expected value for LR: -0.0002947812564940089
Summary Plot for SHAP Values in Test Set:

SHAP Bar Plot for SHAP Values Test Set:

SHAP Decision Plot for SHAP Values in Test Set:

SHAP Decision Plot for Single-Prediction in Test Set:

Checking feature importance for LR2...

Checking for list of csv files...

```
['/Users/jessicakim/Desktop/STREAMLINE/DemoData/Output/hcc_demo/hcc-data_example_no_covariates/model_evaluation/shap_v
alues/testResults/LR_2_shapFIVValues_Test.csv']
```

Unnamed: 0	Features	Importance
0	Performance Status*	1.800500e-04
1	Liver Metastasis	1.477336e-04
2	Alkaline phosphatase (U/L)	1.303614e-04
3	Ferritin (ng/mL)	1.248654e-04
4	Portal Vein Thrombosis	1.208609e-04
5	Albumin (mg/dL)	1.207036e-04
6	Haemoglobin (g/dL)	1.101130e-04
7	Ascites degree*	1.076040e-04
8	Leukocytes(G/L)	1.041162e-04
9	Number of Nodules	9.534978e-05
10	Major dimension of nodule (cm)	9.417302e-05
11	Iron	8.846587e-05
12	Diabetes	8.574631e-05
13	Direct Bilirubin (mg/dL)	8.389784e-05
14	Aspartate transaminase (U/L)	7.283173e-05
15	Platelets	7.035087e-05
16	Gamma glutamyl transferase (U/L)	6.421909e-05
17	Arterial Hypertension	6.210794e-05
18	International Normalised Ratio*	5.933785e-05
19	Human Immunodeficiency Virus	4.465375e-05
20	Portal Hypertension	4.330418e-05
21	Smoking	4.268261e-05
22	Hepatitis B Surface Antigen	3.748354e-05
23	Alcohol	3.611043e-05
24	Mean Corpuscular Volume	3.431638e-05
25	Hepatitis C Virus Antibody	2.915717e-05
26	Chronic Renal Insufficiency	2.643969e-05
27	Cirrhosis	2.100546e-05
28	Hepatitis B e Antigen	1.625872e-05
29	Hemochromatosis	1.581372e-05
30	Creatinine (mg/dL)	1.568307e-05

31	31	Nonalcoholic Steatohepatitis	8.756145e-06
32	32	Packs of cigarets per year	6.604622e-06
33	33	Oxygen Saturation (%)	3.884897e-06
34	34	Endemic Countries	3.504321e-06
35	35	Alpha-Fetoprotein (ng/mL)	4.210016e-07

Out [18]:

Unnamed: 0		Features	Importance
0	0	Performance Status*	1.800500e-04
1	1	Liver Metastasis	1.477336e-04
2	2	Alkaline phosphatase (U/L)	1.303614e-04
3	3	Ferritin (ng/mL)	1.248654e-04
4	4	Portal Vein Thrombosis	1.208609e-04
5	5	Albumin (mg/dL)	1.207036e-04
6	6	Haemoglobin (g/dL)	1.101130e-04
7	7	Ascites degree*	1.076040e-04
8	8	Leukocytes(G/L)	1.041162e-04
9	9	Number of Nodules	9.534978e-05
10	10	Major dimension of nodule (cm)	9.417302e-05
11	11	Iron	8.846587e-05
12	12	Diabetes	8.574631e-05
13	13	Direct Bilirubin (mg/dL)	8.389784e-05
14	14	Aspartate transaminase (U/L)	7.283173e-05
15	15	Platelets	7.035087e-05
16	16	Gamma glutamyl transferase (U/L)	6.421909e-05
17	17	Arterial Hypertension	6.210794e-05
18	18	International Normalised Ratio*	5.933785e-05
19	19	Human Immunodeficiency Virus	4.465375e-05
20	20	Portal Hypertension	4.330418e-05
21	21	Smoking	4.268261e-05
22	22	Hepatitis B Surface Antigen	3.748354e-05
23	23	Alcohol	3.611043e-05
24	24	Mean Corpuscular Volume	3.431638e-05
25	25	Hepatitis C Virus Antibody	2.915717e-05
26	26	Chronic Renal Insufficiency	2.643969e-05
27	27	Cirrhosis	2.100546e-05
28	28	Hepatitis B e Antigen	1.625872e-05
29	29	Hemochromatosis	1.581372e-05
30	30	Creatinine (mg/dL)	1.568307e-05
31	31	Nonalcoholic Steatohepatitis	8.756145e-06
32	32	Packs of cigarets per year	6.604622e-06
33	33	Oxygen Saturation (%)	3.884897e-06
34	34	Endemic Countries	3.504321e-06
35	35	Alpha-Fetoprotein (ng/mL)	4.210016e-07

DT
DT0 In CV0...

Checking if correct model is loaded...
DecisionTreeClassifier(max_depth=17, min_samples_leaf=35, min_samples_split=45, random_state=42)
['Alanine transaminase (U/L)', 'Albumin (mg/dL)', 'Alcohol', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/mL)', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Chronic Renal Insufficiency', 'Cirrhosis', 'Creatinine (mg/dL)', 'Diabetes', 'Direct Bilirubin (mg/dL)', 'Encephalopathy degree*', 'Endemic Countries', 'Esophageal Varices', 'Ferritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haemoglobin (g/dL)', 'Hepatitis B Core Antibody', 'Hepatitis B Surface Antigen', 'Hepatitis C Virus Antibody', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Nonalcoholic Steatohepatitis', 'Number of Nodules', 'Oxygen Saturation (%)', 'Packs of cigarettes per year', 'Performance Status*', 'Portal Vein Thrombosis', 'Splenomegaly', 'Symptoms ', 'Total Bilirubin(mg/dL)', 'Total Proteins (g/dL)']
['Alanine transaminase (U/L)', 'Albumin (mg/dL)', 'Alcohol', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/mL)', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Chronic Renal Insufficiency', 'Cirrhosis', 'Creatinine (mg/dL)', 'Diabetes', 'Direct Bilirubin (mg/dL)', 'Encephalopathy degree*', 'Endemic Countries', 'Esophageal Varices', 'Ferritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haemoglobin (g/dL)', 'Hepatitis B Core Antibody', 'Hepatitis B Surface Antigen', 'Hepatitis C Virus Antibody', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Nonalcoholic Steatohepatitis', 'Number of Nodules', 'Oxygen Saturation (%)', 'Packs of cigarettes per year', 'Performance Status*', 'Portal Vein Thrombosis', 'Splenomegaly', 'Symptoms ', 'Total Bilirubin(mg/dL)', 'Total Proteins (g/dL)']

Checking explainer for DT0...
<shap.explainers._tree.Tree object at 0x7f8905dfc460>

Checking shap values for DT0...

```
[array([[0., 0., 0., ..., 0., 0., 0.],
        [0., 0., 0., ..., 0., 0., 0.],
        [0., 0., 0., ..., 0., 0., 0.],
        ...,
        [0., 0., 0., ..., 0., 0., 0.],
        [0., 0., 0., ..., 0., 0., 0.],
        [0., 0., 0., ..., 0., 0., 0.])), array([[0., 0., 0., ..., 0., 0., 0.],
        [0., 0., 0., ..., 0., 0., 0.],
        [0., 0., 0., ..., 0., 0., 0.],
        ...,
        [0., 0., 0., ..., 0., 0., 0.],
        [0., 0., 0., ..., 0., 0., 0.],
        [0., 0., 0., ..., 0., 0., 0.]])]
```

Checking shap plots for DT0...

Expected value for DT: [0.57272727 0.42727273]
Bar Summary Plot for SHAP Values in Class 0 & 1 in Test Set:

Decision Plot for SHAP Values from Class 0 in Test Set:

Decision Plot for SHAP Values from Class 1 in Test Set:

Checking feature importance for DT0...

Checking for list of csv files...

['/Users/jessicakim/Desktop/STREAMLINE/DemoData/Output/hcc_demo/hcc-data_example_no_covariates/model_evaluation/shap_values/testResults/DT_0_shapFIVValues_Test.csv']

Unnamed: 0	Features	Importance
0	1 Alpha-Fetoprotein (ng/mL)	0.450314
1	2 Alanine transaminase (U/L)	0.000000
2	3 Number of Nodules	0.000000
3	4 International Normalised Ratio*	0.000000
4	5 Iron	0.000000
5	6 Leukocytes(G/L)	0.000000
6	7 Liver Metastasis	0.000000
7	8 Major dimension of nodule (cm)	0.000000
8	9 Mean Corpuscular Volume	0.000000
9	10 Nonalcoholic Steatohepatitis	0.000000
10	11 Oxygen Saturation (%)	0.000000
11	12 Hepatitis B Surface Antigen	0.000000
12	13 Packs of cigarettes per year	0.000000
13	14 Performance Status*	0.000000
14	15 Portal Vein Thrombosis	0.000000
15	16 Splenomegaly	0.000000
16	17 Symptoms	0.000000
17	18 Total Bilirubin(mg/dL)	0.000000
18	19 Hepatitis C Virus Antibody	0.000000
19	20 Hepatitis B Core Antibody	0.000000
20	21 Albumin (mg/dL)	0.000000
21	22 Creatinine (mg/dL)	0.000000
22	23 Alcohol	0.000000
23	24 Alkaline phosphatase (U/L)	0.000000
24	25 Ascites degree*	0.000000
25	26 Aspartate transaminase (U/L)	0.000000
26	27 Chronic Renal Insufficiency	0.000000
27	28 Cirrhosis	0.000000
28	29 Diabetes	0.000000
29	30 Haemoglobin (g/dL)	0.000000
30	31 Direct Bilirubin (mg/dL)	0.000000

31	32	Encephalopathy degree*	0.000000
32	33	Endemic Countries	0.000000
33	34	Esophageal Varices	0.000000
34	35	Ferritin (ng/mL)	0.000000
35	36	Gamma glutamyl transferase (U/L)	0.000000
36	37	Total Proteins (g/dL)	0.000000

Out [18]:

Unnamed: 0		Features	Importance
0	1	Alpha-Fetoprotein (ng/mL)	0.450314
1	2	Alanine transaminase (U/L)	0.000000
2	3	Number of Nodules	0.000000
3	4	International Normalised Ratio*	0.000000
4	5	Iron	0.000000
5	6	Leukocytes(G/L)	0.000000
6	7	Liver Metastasis	0.000000
7	8	Major dimension of nodule (cm)	0.000000
8	9	Mean Corpuscular Volume	0.000000
9	10	Nonalcoholic Steatohepatitis	0.000000
10	11	Oxygen Saturation (%)	0.000000
11	12	Hepatitis B Surface Antigen	0.000000
12	13	Packs of cigarets per year	0.000000
13	14	Performance Status*	0.000000
14	15	Portal Vein Thrombosis	0.000000
15	16	Splenomegaly	0.000000
16	17	Symptoms	0.000000
17	18	Total Bilirubin(mg/dL)	0.000000
18	19	Hepatitis C Virus Antibody	0.000000
19	20	Hepatitis B Core Antibody	0.000000
20	21	Albumin (mg/dL)	0.000000
21	22	Creatinine (mg/dL)	0.000000
22	23	Alcohol	0.000000
23	24	Alkaline phosphatase (U/L)	0.000000
24	25	Ascites degree*	0.000000
25	26	Aspartate transaminase (U/L)	0.000000
26	27	Chronic Renal Insufficiency	0.000000
27	28	Cirrhosis	0.000000
28	29	Diabetes	0.000000
29	30	Haemoglobin (g/dL)	0.000000
30	31	Direct Bilirubin (mg/dL)	0.000000
31	32	Encephalopathy degree*	0.000000
32	33	Endemic Countries	0.000000
33	34	Esophageal Varices	0.000000
34	35	Ferritin (ng/mL)	0.000000
35	36	Gamma glutamyl transferase (U/L)	0.000000
36	37	Total Proteins (g/dL)	0.000000

DT1 In CV1...

```
Checking if correct model is loaded...
DecisionTreeClassifier(criterion='entropy', max_depth=30, min_samples_leaf=20,
                        min_samples_split=5, random_state=42, splitter='random')
['Alanine transaminase (U/L)', 'Albumin (mg/dL)', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/mL)', 'Arterial Hypertension', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Cirrhosis', 'Creatinine (mg/dL)', 'Diabetes', 'Direct Bilirubin (mg/dL)', 'Encephalopathy degree*', 'Endemic Countries', 'Ferritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haemoglobin (g/dL)', 'Hemochromatosis', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibody', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Nonalcoholic Steatohepatitis', 'Number of Nodules', 'Obesity', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Platelets', 'Portal Hypertension', 'Portal Vein Thrombosis', 'Symptoms ', 'Total Bilirubin(mg/dL)']
['Alanine transaminase (U/L)', 'Albumin (mg/dL)', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/mL)', 'Arterial Hypertension', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Cirrhosis', 'Creatinine (mg/dL)', 'Diabetes', 'Direct Bilirubin (mg/dL)', 'Encephalopathy degree*', 'Endemic Countries', 'Ferritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haemoglobin (g/dL)', 'Hemochromatosis', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibody', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Nonalcoholic Steatohepatitis', 'Number of Nodules', 'Obesity', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Platelets', 'Portal Hypertension', 'Portal Vein Thrombosis', 'Symptoms ', 'Total Bilirubin(mg/dL)']
```

Checking explainer for DT1...
<shap.explainers._tree.Tree object at 0x7f88a46b73a0>

Checking shap values for DT1...

```
[array([[ 0.          ,  0.08222611,  0.          , ...,  0.          ,
         0.          ,  0.          ],
       [ 0.          , -0.02219251,  0.          , ...,  0.          ,
         0.          ,  0.          ],
       [ 0.          , -0.02219251,  0.          , ...,  0.          ,
         0.          ,  0.          ],
       ...,
       [ 0.          , -0.02219251,  0.          , ...,  0.          ,
         0.          ,  0.          ],
       [ 0.          , -0.06287879,  0.          , ...,  0.          ,
         0.          ,  0.          ],
       [ 0.          ,  0.02902098,  0.          , ...,  0.          ,
         0.          ,  0.          ]]), array([[ 0.          , -0.08222611,  0.          , ...,  0.          ,
         0.          ,  0.          ],
       [ 0.          ,  0.02219251,  0.          , ...,  0.          ,
         0.          ,  0.          ],
       [ 0.          ,  0.02219251,  0.          , ...,  0.          ,
         0.          ,  0.          ],
       ...,
       [ 0.          ,  0.02219251,  0.          , ...,  0.          ,
         0.          ,  0.          ],
       [ 0.          ,  0.06287879,  0.          , ...,  0.          ,
         0.          ,  0.          ],
       [ 0.          , -0.02902098,  0.          , ...,  0.          ,
         0.          ,  0.          ]])]
```

Checking shap plots for DT1...

Expected value for DT: [0.63636364 0.36363636]
Bar Summary Plot for SHAP Values in Class 0 & 1 in Test Set:

Decision Plot for SHAP Values from Class 0 in Test Set:

Decision Plot for SHAP Values from Class 1 in Test Set:

Checking feature importance for DT1...

Checking for list of csv files...

```
['/Users/jessicakim/Desktop/STREAMLINE/DemoData/Output/hcc_demo/hcc-data_example_no_covariates/model_evaluation/shap_v
values/testResults/DT_1_shapFIValues_Test.csv']
```

Unnamed: 0	Features	Importance	
0	1	Haemoglobin (g/dL)	0.340712
1	2	Ascites degree*	0.127908
2	3	Albumin (mg/dL)	0.097006
3	4	Obesity	0.000000
4	5	Leukocytes(G/L)	0.000000
5	6	Liver Metastasis	0.000000
6	7	Major dimension of nodule (cm)	0.000000
7	8	Mean Corpuscular Volume	0.000000
8	9	Nonalcoholic Steatohepatitis	0.000000
9	10	Number of Nodules	0.000000
10	11	Alanine transaminase (U/L)	0.000000
11	12	Iron	0.000000
12	13	Packs of cigarets per year	0.000000
13	14	Performance Status*	0.000000
14	15	Platelets	0.000000
15	16	Portal Hypertension	0.000000
16	17	Portal Vein Thrombosis	0.000000
17	18	Symptoms	0.000000
18	19	Oxygen Saturation (%)	0.000000
19	20	Hepatitis C Virus Antibody	0.000000

20	21	International Normalised Ratio*	0.000000
21	22	Hepatitis B e Antigen	0.000000
22	23	Hemochromatosis	0.000000
23	24	Gamma glutamyl transferase (U/L)	0.000000
24	25	Ferritin (ng/mL)	0.000000
25	26	Endemic Countries	0.000000
26	27	Encephalopathy degree*	0.000000
27	28	Direct Bilirubin (mg/dL)	0.000000
28	29	Diabetes	0.000000
29	30	Creatinine (mg/dL)	0.000000
30	31	Cirrhosis	0.000000
31	32	Aspartate transaminase (U/L)	0.000000
32	33	Arterial Hypertension	0.000000
33	34	Alpha-Fetoprotein (ng/mL)	0.000000
34	35	Alkaline phosphatase (U/L)	0.000000
35	36	Total Bilirubin(mg/dL)	0.000000

Out [18]:

Unnamed: 0		Features	Importance
0	1	Haemoglobin (g/dL)	0.340712
1	2	Ascites degree*	0.127908
2	3	Albumin (mg/dL)	0.097006
3	4	Obesity	0.000000
4	5	Leukocytes(G/L)	0.000000
5	6	Liver Metastasis	0.000000
6	7	Major dimension of nodule (cm)	0.000000
7	8	Mean Corpuscular Volume	0.000000
8	9	Nonalcoholic Steatohepatitis	0.000000
9	10	Number of Nodules	0.000000
10	11	Alanine transaminase (U/L)	0.000000
11	12	Iron	0.000000
12	13	Packs of cigarets per year	0.000000
13	14	Performance Status*	0.000000
14	15	Platelets	0.000000
15	16	Portal Hypertension	0.000000
16	17	Portal Vein Thrombosis	0.000000
17	18	Symptoms	0.000000
18	19	Oxygen Saturation (%)	0.000000
19	20	Hepatitis C Virus Antibody	0.000000
20	21	International Normalised Ratio*	0.000000
21	22	Hepatitis B e Antigen	0.000000
22	23	Hemochromatosis	0.000000
23	24	Gamma glutamyl transferase (U/L)	0.000000
24	25	Ferritin (ng/mL)	0.000000
25	26	Endemic Countries	0.000000
26	27	Encephalopathy degree*	0.000000
27	28	Direct Bilirubin (mg/dL)	0.000000
28	29	Diabetes	0.000000
29	30	Creatinine (mg/dL)	0.000000
30	31	Cirrhosis	0.000000
31	32	Aspartate transaminase (U/L)	0.000000
32	33	Arterial Hypertension	0.000000
33	34	Alpha-Fetoprotein (ng/mL)	0.000000
34	35	Alkaline phosphatase (U/L)	0.000000
35	36	Total Bilirubin(mg/dL)	0.000000

DT2 In CV2...

```
Checking if correct model is loaded...
DecisionTreeClassifier(class_weight='balanced', max_depth=29,
                        min_samples_leaf=30, min_samples_split=45,
                        random_state=42)
['Albumin (mg/dL)', 'Alcohol', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/mL)', 'Arterial Hypertension', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Chronic Renal Insufficiency', 'Cirrhosis', 'Creatinine (mg/dL)', 'Diabetes', 'Direct Bilirubin (mg/dL)', 'Endemic Countries', 'Ferritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haemoglobin (g/dL)', 'Hemochromatosis', 'Hepatitis B Surface Antigen', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibody', 'Human Immunodeficiency Virus', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Nonalcoholic Steatohepatitis', 'Number of Nodules', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Platelets', 'Portal Hypertension', 'Portal Vein Thrombosis', 'Smoking']
['Albumin (mg/dL)', 'Alcohol', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/mL)', 'Arterial Hypertension', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Chronic Renal Insufficiency', 'Cirrhosis', 'Creatinine (mg/dL)', 'Diabetes', 'Direct Bilirubin (mg/dL)', 'Endemic Countries', 'Ferritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haemoglobin (g/dL)', 'Hemochromatosis', 'Hepatitis B Surface Antigen', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibody', 'Human Immunodeficiency Virus', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Nonalcoholic Steatohepatitis', 'Number of Nodules', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Platelets', 'Portal Hypertension', 'Portal Vein Thrombosis', 'Smoking']
```

Checking explainer for DT2...
<shap.explainers._tree.Tree object at 0x7f8915bbb910>

Checking shap values for DT2...

```
[array([[ -0.14385676,  0.          ,  0.20558515, ...,  0.          ,
         0.          ,  0.          ],
       [ -0.0462963 ,  0.          , -0.18566247, ...,  0.          ,
         0.          ,  0.          ],
       [ -0.0462963 ,  0.          , -0.18566247, ...,  0.          ,
         0.          ,  0.          ],
       ...,
       [ -0.0462963 ,  0.          , -0.18566247, ...,  0.          ,
         0.          ,  0.          ],
       [  0.05769231,  0.          , -0.28965107, ...,  0.          ,
         0.          ,  0.          ],
       [ -0.14385676,  0.          ,  0.20558515, ...,  0.          ,
         0.          ,  0.          ]]), array([[ 0.14385676,  0.          , -0.20558515, ...,  0.          ,
         0.          ,  0.          ],
       [  0.0462963 ,  0.          ,  0.18566247, ...,  0.          ,
         0.          ,  0.          ],
       [  0.0462963 ,  0.          ,  0.18566247, ...,  0.          ,
         0.          ,  0.          ],
       ...,
       [  0.0462963 ,  0.          ,  0.18566247, ...,  0.          ,
         0.          ,  0.          ],
       [-0.05769231,  0.          ,  0.28965107, ...,  0.          ,
         0.          ,  0.          ],
       [  0.14385676,  0.          , -0.20558515, ...,  0.          ,
         0.          ,  0.          ]]])]
```

Checking shap plots for DT2...
Expected value for DT: [0.5 0.5]
Bar Summary Plot for SHAP Values in Class 0 & 1 in Test Set:

Decision Plot for SHAP Values from Class 0 in Test Set:

Decision Plot for SHAP Values from Class 1 in Test Set:

Checking feature importance for DT2...

Checking for list of csv files...

```
['/Users/jessicakim/Desktop/STREAMLINE/DemoData/Output/hcc_demo/hcc-data_example_no_covariates/model_evaluation/shap_v
alues/testResults/DT_2_shapFIValues_Test.csv']
```

Unnamed: 0	Features	Importance
0	1 Alkaline phosphatase (U/L)	0.455851
1	2 Albumin (mg/dL)	0.204616
2	3 Nonalcoholic Steatohepatitis	0.000000
3	4 International Normalised Ratio*	0.000000
4	5 Iron	0.000000
5	6 Leukocytes(G/L)	0.000000
6	7 Liver Metastasis	0.000000
7	8 Major dimension of nodule (cm)	0.000000
8	9 Mean Corpuscular Volume	0.000000
9	10 Number of Nodules	0.000000
10	11 Hepatitis C Virus Antibody	0.000000
11	12 Oxygen Saturation (%)	0.000000
12	13 Packs of cigarets per year	0.000000
13	14 Performance Status*	0.000000
14	15 Platelets	0.000000
15	16 Portal Hypertension	0.000000
16	17 Portal Vein Thrombosis	0.000000
17	18 Human Immunodeficiency Virus	0.000000
18	19 Hepatitis B e Antigen	0.000000

19	20	Alcohol	0.000000
20	21	Creatinine (mg/dL)	0.000000
21	22	Alpha-Fetoprotein (ng/mL)	0.000000
22	23	Arterial Hypertension	0.000000
23	24	Ascites degree*	0.000000
24	25	Aspartate transaminase (U/L)	0.000000
25	26	Chronic Renal Insufficiency	0.000000
26	27	Cirrhosis	0.000000
27	28	Diabetes	0.000000
28	29	Hepatitis B Surface Antigen	0.000000
29	30	Direct Bilirubin (mg/dL)	0.000000
30	31	Endemic Countries	0.000000
31	32	Ferritin (ng/mL)	0.000000
32	33	Gamma glutamyl transferase (U/L)	0.000000
33	34	Haemoglobin (g/dL)	0.000000
34	35	Hemochromatosis	0.000000
35	36	Smoking	0.000000

Out [18]:

Unnamed: 0		Features	Importance
0	1	Alkaline phosphatase (U/L)	0.455851
1	2	Albumin (mg/dL)	0.204616
2	3	Nonalcoholic Steatohepatitis	0.000000
3	4	International Normalised Ratio*	0.000000
4	5	Iron	0.000000
5	6	Leukocytes(G/L)	0.000000
6	7	Liver Metastasis	0.000000
7	8	Major dimension of nodule (cm)	0.000000
8	9	Mean Corpuscular Volume	0.000000
9	10	Number of Nodules	0.000000
10	11	Hepatitis C Virus Antibody	0.000000
11	12	Oxygen Saturation (%)	0.000000
12	13	Packs of cigarets per year	0.000000
13	14	Performance Status*	0.000000
14	15	Platelets	0.000000
15	16	Portal Hypertension	0.000000
16	17	Portal Vein Thrombosis	0.000000
17	18	Human Immunodeficiency Virus	0.000000
18	19	Hepatitis B e Antigen	0.000000
19	20	Alcohol	0.000000
20	21	Creatinine (mg/dL)	0.000000
21	22	Alpha-Fetoprotein (ng/mL)	0.000000
22	23	Arterial Hypertension	0.000000
23	24	Ascites degree*	0.000000
24	25	Aspartate transaminase (U/L)	0.000000
25	26	Chronic Renal Insufficiency	0.000000
26	27	Cirrhosis	0.000000
27	28	Diabetes	0.000000
28	29	Hepatitis B Surface Antigen	0.000000
29	30	Direct Bilirubin (mg/dL)	0.000000
30	31	Endemic Countries	0.000000
31	32	Ferritin (ng/mL)	0.000000
32	33	Gamma glutamyl transferase (U/L)	0.000000
33	34	Haemoglobin (g/dL)	0.000000
34	35	Hemochromatosis	0.000000
35	36	Smoking	0.000000

```
RF
RF0 In CV0...

Checking if correct model is loaded...
RandomForestClassifier(max_depth=9, max_features=None, min_samples_leaf=9,
                        min_samples_split=24, n_estimators=935, random_state=42)
['Alanine transaminase (U/L)', 'Albumin (mg/dL)', 'Alcohol', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/m
L)', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Chronic Renal Insufficiency', 'Cirrhosis', 'Creatinine (mg/d
L)', 'Diabetes', 'Direct Bilirubin (mg/dL)', 'Encephalopathy degree*', 'Endemic Countries', 'Esophageal Varices', 'Fer
ritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haemoglobin (g/dL)', 'Hepatitis B Core Antibody', 'Hepatitis B Su
rface Antigen', 'Hepatitis C Virus Antibody', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Met
astasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Nonalcoholic Steatohepatitis', 'Number of Nodu
les', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Portal Vein Thrombosis', 'Splenom
egaly', 'Symptoms ', 'Total Bilirubin(mg/dL)', 'Total Proteins (g/dL)']
['Alanine transaminase (U/L)', 'Albumin (mg/dL)', 'Alcohol', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/m
L)', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Chronic Renal Insufficiency', 'Cirrhosis', 'Creatinine (mg/d
L)', 'Diabetes', 'Direct Bilirubin (mg/dL)', 'Encephalopathy degree*', 'Endemic Countries', 'Esophageal Varices', 'Fer
ritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haemoglobin (g/dL)', 'Hepatitis B Core Antibody', 'Hepatitis B Su
rface Antigen', 'Hepatitis C Virus Antibody', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Met
astasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Nonalcoholic Steatohepatitis', 'Number of Nodu
les', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Portal Vein Thrombosis', 'Splenom
egaly', 'Symptoms ', 'Total Bilirubin(mg/dL)', 'Total Proteins (g/dL)']

Checking explainer for RF0...
<shap.explainers._tree.Tree object at 0x7f8905deefa0>

Checking shap values for RF0...

[array([[ -8.04892952e-04,  3.49123556e-02, -7.39028100e-04, ...,
        -2.55865722e-05, -4.80373900e-04, -5.51427280e-03],
       [ 8.98581287e-04, -2.36897659e-02, -7.01881775e-05, ...,
        -4.69087156e-05, -1.53604128e-03, -9.93491258e-03],
       [ 1.09297656e-04, -2.10001659e-02,  2.39694450e-04, ...,
        -4.18242045e-05, -3.61466526e-04, -6.98520090e-03],
       ...,
       [ 3.18715672e-04,  2.90540264e-02,  2.67651465e-04, ...,
        -4.18242045e-05,  7.09998438e-05,  7.60365635e-03],
       [-1.87966055e-03, -1.58333798e-02,  2.67651465e-04, ...,
        3.18309142e-04,  1.93293842e-03, -7.31627298e-04],
       [-1.82586614e-03, -2.32531905e-02,  1.25596859e-04, ...,
        3.18309142e-04, -1.88973059e-03, -1.80372814e-03]])], array([[ 8.04892952e-04, -3.49123556e-02,  7.39028100e-0
4, ...,
        2.55865722e-05,  4.80373900e-04,  5.51427280e-03],
       [-8.98581287e-04,  2.36897659e-02,  7.01881775e-05, ...,
        4.69087156e-05,  1.53604128e-03,  9.93491258e-03],
       [-1.09297656e-04,  2.10001659e-02, -2.39694450e-04, ...,
        4.18242045e-05,  3.61466526e-04,  6.98520090e-03],
       ...,
       [-3.18715672e-04, -2.90540264e-02, -2.67651465e-04, ...,
        4.18242045e-05, -7.09998438e-05, -7.60365635e-03],
       [ 1.87966055e-03,  1.58333798e-02, -2.67651465e-04, ...,
        -3.18309142e-04, -1.93293842e-03,  7.31627298e-04],
       [ 1.82586614e-03,  2.32531905e-02, -1.25596859e-04, ...,
        -3.18309142e-04,  1.88973059e-03,  1.80372814e-03]])])

Checking shap plots for RF0...

Expected value for RF: [0.57252309 0.42747691]
Bar Summary Plot for SHAP Values in Class 0 & 1 in Test Set:

Decision Plot for SHAP Values from Class 0 in Test Set:

Decision Plot for SHAP Values from Class 1 in Test Set:

Checking feature importance for RF0...

Checking for list of csv files...

['/Users/jessicakim/Desktop/STREAMLINE/DemoData/Output/hcc_demo/hcc-data_example_no_covariates/model_evaluation/shap_v
alues/testResults/RF_0_shapFIValues_Test.csv']
  Unnamed: 0      Features  Importance
0           1  Alpha-Fetoprotein (ng/mL)  0.298214
1           2                Iron  0.066051
2           3      Ferritin (ng/mL)  0.063013
3           4      Albumin (mg/dL)  0.056744
4           5  Aspartate transaminase (U/L)  0.037689
5           6  Alkaline phosphatase (U/L)  0.036931
6           7      Performance Status*  0.027310
7           8  Major dimension of nodule (cm)  0.024439
8           9      Creatinine (mg/dL)  0.022242
9          10      Haemoglobin (g/dL)  0.021447
10          11  Direct Bilirubin (mg/dL)  0.013847
11          12      Total Proteins (g/dL)  0.013708
12          13      Number of Nodules  0.011870
13          14  International Normalised Ratio*  0.010403
14          15      Oxygen Saturation (%)  0.008767
15          16      Liver Metastasis  0.008502
16          17  Gamma glutamyl transferase (U/L)  0.007045
17          18      Leukocytes(G/L)  0.006241
```

18	19	Mean Corpuscular Volume	0.004768
19	20	Packs of cigarets per year	0.004199
20	21	Alanine transaminase (U/L)	0.003199
21	22	Diabetes	0.002952
22	23	Ascites degree*	0.002070
23	24	Total Bilirubin(mg/dL)	0.002040
24	25	Esophageal Varices	0.001880
25	26	Hepatitis B Core Antibody	0.000923
26	27	Portal Vein Thrombosis	0.000551
27	28	Alcohol	0.000485
28	29	Hepatitis C Virus Antibody	0.000389
29	30	Symptoms	0.000244
30	31	Chronic Renal Insufficiency	0.000195
31	32	Splenomegaly	0.000172
32	33	Hepatitis B Surface Antigen	0.000000
33	34	Endemic Countries	0.000000
34	35	Encephalopathy degree*	0.000000
35	36	Nonalcoholic Steatohepatitis	0.000000
36	37	Cirrhosis	0.000000

Out[18]:

Unnamed: 0		Features	Importance
0	1	Alpha-Fetoprotein (ng/mL)	0.298214
1	2	Iron	0.066051
2	3	Ferritin (ng/mL)	0.063013
3	4	Albumin (mg/dL)	0.056744
4	5	Aspartate transaminase (U/L)	0.037689
5	6	Alkaline phosphatase (U/L)	0.036931
6	7	Performance Status*	0.027310
7	8	Major dimension of nodule (cm)	0.024439
8	9	Creatinine (mg/dL)	0.022242
9	10	Haemoglobin (g/dL)	0.021447
10	11	Direct Bilirubin (mg/dL)	0.013847
11	12	Total Proteins (g/dL)	0.013708
12	13	Number of Nodules	0.011870
13	14	International Normalised Ratio*	0.010403
14	15	Oxygen Saturation (%)	0.008767
15	16	Liver Metastasis	0.008502
16	17	Gamma glutamyl transferase (U/L)	0.007045
17	18	Leukocytes(G/L)	0.006241
18	19	Mean Corpuscular Volume	0.004768
19	20	Packs of cigarets per year	0.004199
20	21	Alanine transaminase (U/L)	0.003199
21	22	Diabetes	0.002952
22	23	Ascites degree*	0.002070
23	24	Total Bilirubin(mg/dL)	0.002040
24	25	Esophageal Varices	0.001880
25	26	Hepatitis B Core Antibody	0.000923
26	27	Portal Vein Thrombosis	0.000551
27	28	Alcohol	0.000485
28	29	Hepatitis C Virus Antibody	0.000389
29	30	Symptoms	0.000244
30	31	Chronic Renal Insufficiency	0.000195
31	32	Splenomegaly	0.000172
32	33	Hepatitis B Surface Antigen	0.000000
33	34	Endemic Countries	0.000000
34	35	Encephalopathy degree*	0.000000
35	36	Nonalcoholic Steatohepatitis	0.000000
36	37	Cirrhosis	0.000000

RF1 In CV1...

```
Checking if correct model is loaded...
RandomForestClassifier(class_weight='balanced', max_depth=25,
                        max_features='log2', min_samples_leaf=16,
                        min_samples_split=45, n_estimators=194, oob_score=True,
                        random_state=42)
['Alanine transaminase (U/L)', 'Albumin (mg/dL)', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/mL)', 'Arterial Hypertension', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Cirrhosis', 'Creatinine (mg/dL)', 'Diabetes', 'Direct Bilirubin (mg/dL)', 'Encephalopathy degree*', 'Endemic Countries', 'Ferritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haemoglobin (g/dL)', 'Hemochromatosis', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibody', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Nonalcoholic Steatohepatitis', 'Number of Nodules', 'Obesity', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Platelets', 'Portal Hypertension', 'Portal Vein Thrombosis', 'Symptoms ', 'Total Bilirubin(mg/dL)']
['Alanine transaminase (U/L)', 'Albumin (mg/dL)', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/mL)', 'Arterial Hypertension', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Cirrhosis', 'Creatinine (mg/dL)', 'Diabetes', 'Direct Bilirubin (mg/dL)', 'Encephalopathy degree*', 'Endemic Countries', 'Ferritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haemoglobin (g/dL)', 'Hemochromatosis', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibody', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Nonalcoholic Steatohepatitis', 'Number of Nodules', 'Obesity', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Platelets', 'Portal Hypertension', 'Portal Vein Thrombosis', 'Symptoms ', 'Total Bilirubin(mg/dL)']
```

Checking explainer for RF1...
<shap.explainers._tree.Tree object at 0x7f8905dafa90>

Checking shap values for RF1...

```
[array([[ 6.89282193e-04,  5.16384335e-02, -1.76947858e-02, ...,
         4.15186605e-04, -5.22220942e-03,  1.52801529e-02],
        [-2.86516996e-03, -1.75170521e-03,  1.53660207e-02, ...,
         1.57204645e-04,  1.14384010e-02,  1.36293862e-02],
        [ 9.58343029e-04, -3.08481544e-02, -1.63920514e-02, ...,
         1.57204645e-04,  1.17329978e-02, -6.96607661e-03],
        ...,
        [-4.62392460e-05,  2.51854494e-02, -1.70193937e-02, ...,
         1.57204645e-04,  1.17917953e-02,  1.21731827e-02],
        [ 8.00804127e-04, -3.08295224e-02, -1.58989779e-02, ...,
        -1.04418188e-03, -5.37882055e-03, -6.99630666e-04],
        [-1.58770860e-03,  5.12094387e-02, -9.49257495e-03, ...,
         1.57204645e-04, -5.09113830e-03,  1.43028838e-02]]), array([[ -6.89282193e-04, -5.16384335e-02,  1.76947858e-0
2, ...,
        -4.15186605e-04,  5.22220942e-03, -1.52801529e-02],
        [ 2.86516996e-03,  1.75170521e-03, -1.53660207e-02, ...,
        -1.57204645e-04, -1.14384010e-02, -1.36293862e-02],
        [-9.58343029e-04,  3.08481544e-02,  1.63920514e-02, ...,
        -1.57204645e-04, -1.17329978e-02,  6.96607661e-03],
        ...,
        [ 4.62392460e-05, -2.51854494e-02,  1.70193937e-02, ...,
        -1.57204645e-04, -1.17917953e-02, -1.21731827e-02],
        [-8.00804127e-04,  3.08295224e-02,  1.58989779e-02, ...,
         1.04418188e-03,  5.37882055e-03,  6.99630666e-04],
        [ 1.58770860e-03, -5.12094387e-02,  9.49257495e-03, ...,
        -1.57204645e-04,  5.09113830e-03, -1.43028838e-02]])]
```

Checking shap plots for RF1...

Expected value for RF: [0.49611628 0.50388372]
Bar Summary Plot for SHAP Values in Class 0 & 1 in Test Set:

Decision Plot for SHAP Values from Class 0 in Test Set:

Decision Plot for SHAP Values from Class 1 in Test Set:

Checking feature importance for RF1...

Checking for list of csv files...

```
['/Users/jessicakim/Desktop/STREAMLINE/DemoData/Output/hcc_demo/hcc-data_example_no_covariates/model_evaluation/shap_v
alues/testResults/RF_1_shapFIValues_Test.csv']
```

Unnamed: 0	Features	Importance
0	1 Albumin (mg/dL)	0.068225
1	2 Haemoglobin (g/dL)	0.065448
2	3 Iron	0.043139
3	4 Ascites degree*	0.034337
4	5 Alpha-Fetoprotein (ng/mL)	0.032638
5	6 Performance Status*	0.032205
6	7 Direct Bilirubin (mg/dL)	0.029600
7	8 Total Bilirubin(mg/dL)	0.027581
8	9 Oxygen Saturation (%)	0.027019
9	10 Alkaline phosphatase (U/L)	0.026121
10	11 International Normalised Ratio*	0.017099
11	12 Symptoms	0.015247
12	13 Major dimension of nodule (cm)	0.010214
13	14 Mean Corpuscular Volume	0.007092
14	15 Ferritin (ng/mL)	0.006220
15	16 Aspartate transaminase (U/L)	0.005399
16	17 Gamma glutamyl transferase (U/L)	0.004433

17	18	Number of Nodules	0.003951
18	19	Platelets	0.002901
19	20	Alanine transaminase (U/L)	0.002648
20	21	Leukocytes(G/L)	0.002368
21	22	Packs of cigarets per year	0.001856
22	23	Portal Hypertension	0.001280
23	24	Diabetes	0.001178
24	25	Portal Vein Thrombosis	0.000723
25	26	Creatinine (mg/dL)	0.000519
26	27	Hepatitis B e Antigen	0.000000
27	28	Liver Metastasis	0.000000
28	29	Hemochromatosis	0.000000
29	30	Endemic Countries	0.000000
30	31	Nonalcoholic Steatohepatitis	0.000000
31	32	Encephalopathy degree*	0.000000
32	33	Obesity	0.000000
33	34	Cirrhosis	0.000000
34	35	Arterial Hypertension	0.000000
35	36	Hepatitis C Virus Antibody	0.000000

Out[18]:

Unnamed: 0		Features	Importance
0	1	Albumin (mg/dL)	0.068225
1	2	Haemoglobin (g/dL)	0.065448
2	3	Iron	0.043139
3	4	Ascites degree*	0.034337
4	5	Alpha-Fetoprotein (ng/mL)	0.032638
5	6	Performance Status*	0.032205
6	7	Direct Bilirubin (mg/dL)	0.029600
7	8	Total Bilirubin(mg/dL)	0.027581
8	9	Oxygen Saturation (%)	0.027019
9	10	Alkaline phosphatase (U/L)	0.026121
10	11	International Normalised Ratio*	0.017099
11	12	Symptoms	0.015247
12	13	Major dimension of nodule (cm)	0.010214
13	14	Mean Corpuscular Volume	0.007092
14	15	Ferritin (ng/mL)	0.006220
15	16	Aspartate transaminase (U/L)	0.005399
16	17	Gamma glutamyl transferase (U/L)	0.004433
17	18	Number of Nodules	0.003951
18	19	Platelets	0.002901
19	20	Alanine transaminase (U/L)	0.002648
20	21	Leukocytes(G/L)	0.002368
21	22	Packs of cigarets per year	0.001856
22	23	Portal Hypertension	0.001280
23	24	Diabetes	0.001178
24	25	Portal Vein Thrombosis	0.000723
25	26	Creatinine (mg/dL)	0.000519
26	27	Hepatitis B e Antigen	0.000000
27	28	Liver Metastasis	0.000000
28	29	Hemochromatosis	0.000000
29	30	Endemic Countries	0.000000
30	31	Nonalcoholic Steatohepatitis	0.000000
31	32	Encephalopathy degree*	0.000000
32	33	Obesity	0.000000
33	34	Cirrhosis	0.000000
34	35	Arterial Hypertension	0.000000
35	36	Hepatitis C Virus Antibody	0.000000

RF2 In CV2...

```
Checking if correct model is loaded...
RandomForestClassifier(class_weight='balanced', max_depth=16, max_features=None,
                        min_samples_leaf=20, min_samples_split=40,
                        n_estimators=299, oob_score=True, random_state=42)
['Albumin (mg/dL)', 'Alcohol', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/mL)', 'Arterial Hypertension', 'As
cites degree*', 'Aspartate transaminase (U/L)', 'Chronic Renal Insufficiency', 'Cirrhosis', 'Creatinine (mg/dL)', 'Dia
betes', 'Direct Bilirubin (mg/dL)', 'Endemic Countries', 'Ferritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haem
oglobin (g/dL)', 'Hemochromatosis', 'Hepatitis B Surface Antigen', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibod
y', 'Human Immunodeficiency Virus', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Metastasis',
'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Nonalcoholic Steatohepatitis', 'Number of Nodules', 'Oxy
gen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Platelets', 'Portal Hypertension', 'Portal
Vein Thrombosis', 'Smoking']
['Albumin (mg/dL)', 'Alcohol', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/mL)', 'Arterial Hypertension', 'As
cites degree*', 'Aspartate transaminase (U/L)', 'Chronic Renal Insufficiency', 'Cirrhosis', 'Creatinine (mg/dL)', 'Dia
betes', 'Direct Bilirubin (mg/dL)', 'Endemic Countries', 'Ferritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haem
oglobin (g/dL)', 'Hemochromatosis', 'Hepatitis B Surface Antigen', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibod
y', 'Human Immunodeficiency Virus', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Metastasis',
'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Nonalcoholic Steatohepatitis', 'Number of Nodules', 'Oxy
gen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Platelets', 'Portal Hypertension', 'Portal
Vein Thrombosis', 'Smoking']
```

Checking explainer for RF2...
<shap.explainers._tree.Tree object at 0x7f88caf8b1c0>

Checking shap values for RF2...

```
[array([[ -1.64930049e-02,  0.00000000e+00,  9.65987236e-02, ...,
          0.00000000e+00,  0.00000000e+00, -2.92612124e-04],
        [ -1.10170786e-02,  0.00000000e+00, -1.61874431e-01, ...,
          0.00000000e+00,  0.00000000e+00,  4.35547992e-04],
        [ -1.17523110e-02,  0.00000000e+00, -1.57953243e-01, ...,
          0.00000000e+00,  0.00000000e+00, -2.92612124e-04],
        ...,
        [ -1.24810033e-02,  0.00000000e+00, -1.55756997e-01, ...,
          0.00000000e+00,  0.00000000e+00,  4.35547992e-04],
        [  2.57852085e-02,  0.00000000e+00, -8.08842985e-02, ...,
          0.00000000e+00,  0.00000000e+00,  4.35547992e-04],
        [ -1.64930049e-02,  0.00000000e+00,  1.95650937e-01, ...,
          0.00000000e+00,  0.00000000e+00, -1.15877556e-04]]), array([[ 1.64930049e-02,  0.00000000e+00, -9.65987236e-0
2, ...,
          0.00000000e+00,  0.00000000e+00,  2.92612124e-04],
        [  1.10170786e-02,  0.00000000e+00,  1.61874431e-01, ...,
          0.00000000e+00,  0.00000000e+00, -4.35547992e-04],
        [  1.17523110e-02,  0.00000000e+00,  1.57953243e-01, ...,
          0.00000000e+00,  0.00000000e+00,  2.92612124e-04],
        ...,
        [  1.24810033e-02,  0.00000000e+00,  1.55756997e-01, ...,
          0.00000000e+00,  0.00000000e+00, -4.35547992e-04],
        [ -2.57852085e-02,  0.00000000e+00,  8.08842985e-02, ...,
          0.00000000e+00,  0.00000000e+00, -4.35547992e-04],
        [  1.64930049e-02,  0.00000000e+00, -1.95650937e-01, ...,
          0.00000000e+00,  0.00000000e+00,  1.15877556e-04]])]
```

Checking shap plots for RF2...

Expected value for RF: [0.49969365 0.50030635]
Bar Summary Plot for SHAP Values in Class 0 & 1 in Test Set:

Decision Plot for SHAP Values from Class 0 in Test Set:

Decision Plot for SHAP Values from Class 1 in Test Set:

Checking feature importance for RF2...

Checking for list of csv files...

```
['/Users/jessicakim/Desktop/STREAMLINE/DemoData/Output/hcc_demo/hcc-data_example_no_covariates/model_evaluation/shap_v
alues/testResults/RF_2_shapFIValues_Test.csv']
```

Unnamed: 0	Features	Importance
0	1 Alkaline phosphatase (U/L)	0.315891
1	2 Ferritin (ng/mL)	0.048037
2	3 Iron	0.046382
3	4 Alpha-Fetoprotein (ng/mL)	0.046257
4	5 Haemoglobin (g/dL)	0.035694
5	6 Albumin (mg/dL)	0.034086
6	7 Leukocytes(G/L)	0.023832
7	8 Performance Status*	0.018199
8	9 Major dimension of nodule (cm)	0.005627
9	10 Platelets	0.004820
10	11 Gamma glutamyl transferase (U/L)	0.003898
11	12 Aspartate transaminase (U/L)	0.003235
12	13 Oxygen Saturation (%)	0.003094
13	14 Creatinine (mg/dL)	0.002749
14	15 Packs of cigarets per year	0.001782
15	16 Number of Nodules	0.001432
16	17 Arterial Hypertension	0.000879
17	18 Mean Corpuscular Volume	0.000830

18	19	Direct Bilirubin (mg/dL)	0.000815
19	20	Diabetes	0.000685
20	21	Smoking	0.000518
21	22	International Normalised Ratio*	0.000000
22	23	Human Immunodeficiency Virus	0.000000
23	24	Hepatitis C Virus Antibody	0.000000
24	25	Liver Metastasis	0.000000
25	26	Alcohol	0.000000
26	27	Hepatitis B Surface Antigen	0.000000
27	28	Nonalcoholic Steatohepatitis	0.000000
28	29	Hemochromatosis	0.000000
29	30	Endemic Countries	0.000000
30	31	Cirrhosis	0.000000
31	32	Chronic Renal Insufficiency	0.000000
32	33	Ascites degree*	0.000000
33	34	Portal Hypertension	0.000000
34	35	Portal Vein Thrombosis	0.000000
35	36	Hepatitis B e Antigen	0.000000

Out[18]:

Unnamed: 0		Features	Importance
0	1	Alkaline phosphatase (U/L)	0.315891
1	2	Ferritin (ng/mL)	0.048037
2	3	Iron	0.046382
3	4	Alpha-Fetoprotein (ng/mL)	0.046257
4	5	Haemoglobin (g/dL)	0.035694
5	6	Albumin (mg/dL)	0.034086
6	7	Leukocytes(G/L)	0.023832
7	8	Performance Status*	0.018199
8	9	Major dimension of nodule (cm)	0.005627
9	10	Platelets	0.004820
10	11	Gamma glutamyl transferase (U/L)	0.003898
11	12	Aspartate transaminase (U/L)	0.003235
12	13	Oxygen Saturation (%)	0.003094
13	14	Creatinine (mg/dL)	0.002749
14	15	Packs of cigarets per year	0.001782
15	16	Number of Nodules	0.001432
16	17	Arterial Hypertension	0.000879
17	18	Mean Corpuscular Volume	0.000830
18	19	Direct Bilirubin (mg/dL)	0.000815
19	20	Diabetes	0.000685
20	21	Smoking	0.000518
21	22	International Normalised Ratio*	0.000000
22	23	Human Immunodeficiency Virus	0.000000
23	24	Hepatitis C Virus Antibody	0.000000
24	25	Liver Metastasis	0.000000
25	26	Alcohol	0.000000
26	27	Hepatitis B Surface Antigen	0.000000
27	28	Nonalcoholic Steatohepatitis	0.000000
28	29	Hemochromatosis	0.000000
29	30	Endemic Countries	0.000000
30	31	Cirrhosis	0.000000
31	32	Chronic Renal Insufficiency	0.000000
32	33	Ascites degree*	0.000000
33	34	Portal Hypertension	0.000000
34	35	Portal Vein Thrombosis	0.000000
35	36	Hepatitis B e Antigen	0.000000


```
XGB
XGB0 In CV0...

Checking if correct model is loaded...
XGBClassifier(alpha=0.0003085901759707382, base_score=0.5, booster='gbtree',
              callbacks=None, colsample_bylevel=1, colsample_bynode=1,
              colsample_bytree=0.31595586732894876, early_stopping_rounds=None,
              enable_categorical=False, eta=0.0016131413768891527,
              eval_metric=None, gamma=1.086786493948363e-07, gpu_id=-1,
              grow_policy='lossguide', importance_type=None,
              interaction_constraints='', learning_rate=0.00161314139,
              max_bin=256, max_cat_to_onehot=4, max_delta_step=0, max_depth=3,
              max_leaves=0, min_child_weight=9.912142174935715,
              min_samples_leaf=32, min_samples_split=43, missing=nan,
              monotone_constraints=('', n_estimators=305, n_jobs=1, nthread=1, ...))
['Alanine transaminase (U/L)', 'Albumin (mg/dL)', 'Alcohol', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/m
L)', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Chronic Renal Insufficiency', 'Cirrhosis', 'Creatinine (mg/d
L)', 'Diabetes', 'Direct Bilirubin (mg/dL)', 'Encephalopathy degree*', 'Endemic Countries', 'Esophageal Varices', 'Fer
ritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haemoglobin (g/dL)', 'Hepatitis B Core Antibody', 'Hepatitis B Su
rface Antigen', 'Hepatitis C Virus Antibody', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Met
astasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Nonalcoholic Steatohepatitis', 'Number of Nodu
les', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Portal Vein Thrombosis', 'Splenom
egaly', 'Symptoms ', 'Total Bilirubin(mg/dL)', 'Total Proteins (g/dL)']
['Alanine transaminase (U/L)', 'Albumin (mg/dL)', 'Alcohol', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/m
L)', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Chronic Renal Insufficiency', 'Cirrhosis', 'Creatinine (mg/d
L)', 'Diabetes', 'Direct Bilirubin (mg/dL)', 'Encephalopathy degree*', 'Endemic Countries', 'Esophageal Varices', 'Fer
ritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haemoglobin (g/dL)', 'Hepatitis B Core Antibody', 'Hepatitis B Su
rface Antigen', 'Hepatitis C Virus Antibody', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Met
astasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Nonalcoholic Steatohepatitis', 'Number of Nodu
les', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Portal Vein Thrombosis', 'Splenom
egaly', 'Symptoms ', 'Total Bilirubin(mg/dL)', 'Total Proteins (g/dL)']

Checking explainer for XGB0...
<shap.explainers._tree.Tree object at 0x7f890200e130>

Checking shap values for XGB0...

[[ 0.00033414 -0.00198181  0.          ...  0.          0.
   0.00147086]
 [-0.0001525  0.00104749  0.          ...  0.          0.
   0.00147086]
 [ 0.00033414 -0.00198181  0.          ...  0.          0.
   0.00174883]
 ...
 [ 0.00013163 -0.00198181  0.          ...  0.          0.
  -0.00234208]
 [-0.0001525  -0.00198181  0.          ...  0.          0.
   0.00110088]
 [ 0.0001405   0.00104749  0.          ...  0.          0.
   0.00082496]]

Checking shap plots for XGB0...

Expected value for XGB: 0.00014352845028042793
Summary Plot for SHAP Values in Test Set:

SHAP Bar Plot for SHAP Values Test Set:

SHAP Decision Plot for SHAP Values in Test Set:

SHAP Decision Plot for Single-Prediction in Test Set:

Checking feature importance for XGB0...

Checking for list of csv files...

['/Users/jessicakim/Desktop/STREAMLINE/DemoData/Output/hcc_demo/hcc-data_example_no_covariates/model_evaluation/shap_v
alues/testResults/XGB_0_shapFIValues_Test.csv']
  Unnamed: 0      Features  Importance
0           0  Alpha-Fetoprotein (ng/mL)  0.092905
1           1  Alkaline phosphatase (U/L)  0.062978
2           2    Performance Status*  0.043804
3           3             Iron  0.037759
4           4  Major dimension of nodule (cm)  0.024830
5           5  Aspartate transaminase (U/L)  0.011059
6           6    Haemoglobin (g/dL)  0.010062
7           7    Creatinine (mg/dL)  0.007855
8           8  Gamma glutamyl transferase (U/L)  0.005661
9           9    Ferritin (ng/mL)  0.004340
10          10    Total Proteins (g/dL)  0.001796
11          11    Direct Bilirubin (mg/dL)  0.001578
12          12      Albumin (mg/dL)  0.001540
13          13  Packs of cigarets per year  0.001346
14          14    Leukocytes(G/L)  0.000883
15          15    Oxygen Saturation (%)  0.000377
16          16  International Normalised Ratio*  0.000336
17          17    Alanine transaminase (U/L)  0.000173
18          18    Total Bilirubin(mg/dL)  0.000000
19          19      Cirrhosis  0.000000
20          20      Symptoms  0.000000
21          21      Alcohol  0.000000
```

22	22	Splenomegaly	0.000000
23	23	Portal Vein Thrombosis	0.000000
24	24	Ascites degree*	0.000000
25	25	Number of Nodules	0.000000
26	26	Chronic Renal Insufficiency	0.000000
27	27	Nonalcoholic Steatohepatitis	0.000000
28	28	Hepatitis B Surface Antigen	0.000000
29	29	Diabetes	0.000000
30	30	Mean Corpuscular Volume	0.000000
31	31	Encephalopathy degree*	0.000000
32	32	Endemic Countries	0.000000
33	33	Esophageal Varices	0.000000
34	34	Liver Metastasis	0.000000
35	35	Hepatitis C Virus Antibody	0.000000
36	36	Hepatitis B Core Antibody	0.000000

Out [18]:

Unnamed: 0		Features	Importance
0	0	Alpha-Fetoprotein (ng/mL)	0.092905
1	1	Alkaline phosphatase (U/L)	0.062978
2	2	Performance Status*	0.043804
3	3	Iron	0.037759
4	4	Major dimension of nodule (cm)	0.024830
5	5	Aspartate transaminase (U/L)	0.011059
6	6	Haemoglobin (g/dL)	0.010062
7	7	Creatinine (mg/dL)	0.007855
8	8	Gamma glutamyl transferase (U/L)	0.005661
9	9	Ferritin (ng/mL)	0.004340
10	10	Total Proteins (g/dL)	0.001796
11	11	Direct Bilirubin (mg/dL)	0.001578
12	12	Albumin (mg/dL)	0.001540
13	13	Packs of cigarets per year	0.001346
14	14	Leukocytes(G/L)	0.000883
15	15	Oxygen Saturation (%)	0.000377
16	16	International Normalised Ratio*	0.000336
17	17	Alanine transaminase (U/L)	0.000173
18	18	Total Bilirubin(mg/dL)	0.000000
19	19	Cirrhosis	0.000000
20	20	Symptoms	0.000000
21	21	Alcohol	0.000000
22	22	Splenomegaly	0.000000
23	23	Portal Vein Thrombosis	0.000000
24	24	Ascites degree*	0.000000
25	25	Number of Nodules	0.000000
26	26	Chronic Renal Insufficiency	0.000000
27	27	Nonalcoholic Steatohepatitis	0.000000
28	28	Hepatitis B Surface Antigen	0.000000
29	29	Diabetes	0.000000
30	30	Mean Corpuscular Volume	0.000000
31	31	Encephalopathy degree*	0.000000
32	32	Endemic Countries	0.000000
33	33	Esophageal Varices	0.000000
34	34	Liver Metastasis	0.000000
35	35	Hepatitis C Virus Antibody	0.000000
36	36	Hepatitis B Core Antibody	0.000000

XGB1 In CV1...

```
Checking if correct model is loaded...
XGBClassifier(alpha=0.2551087479356069, base_score=0.5, booster='gbtree',
              callbacks=None, colsample_bylevel=1, colsample_bynode=1,
              colsample_bytree=0.7752104293797693, early_stopping_rounds=None,
              enable_categorical=False, eta=0.01045236594154273,
              eval_metric=None, gamma=1.2055493072793457e-05, gpu_id=-1,
              grow_policy='depthwise', importance_type=None,
              interaction_constraints='', learning_rate=0.0104523655,
              max_bin=256, max_cat_to_onehot=4, max_delta_step=0, max_depth=11,
              max_leaves=0, min_child_weight=1.1403861951454617,
              min_samples_leaf=23, min_samples_split=24, missing=nan,
              monotone_constraints=('', n_estimators=372, n_jobs=1, nthread=1, ...))
['Alanine transaminase (U/L)', 'Albumin (mg/dL)', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/mL)', 'Arterial Hypertension', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Cirrhosis', 'Creatinine (mg/dL)', 'Diabetes', 'Direct Bilirubin (mg/dL)', 'Encephalopathy degree*', 'Endemic Countries', 'Ferritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haemoglobin (g/dL)', 'Hemochromatosis', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibody', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Nonalcoholic Steatohepatitis', 'Number of Nodules', 'Obesity', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Platelets', 'Portal Hypertension', 'Portal Vein Thrombosis', 'Symptoms ', 'Total Bilirubin(mg/dL)']
['Alanine transaminase (U/L)', 'Albumin (mg/dL)', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/mL)', 'Arterial Hypertension', 'Ascites degree*', 'Aspartate transaminase (U/L)', 'Cirrhosis', 'Creatinine (mg/dL)', 'Diabetes', 'Direct Bilirubin (mg/dL)', 'Encephalopathy degree*', 'Endemic Countries', 'Ferritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haemoglobin (g/dL)', 'Hemochromatosis', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibody', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Metastasis', 'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Nonalcoholic Steatohepatitis', 'Number of Nodules', 'Obesity', 'Oxygen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Platelets', 'Portal Hypertension', 'Portal Vein Thrombosis', 'Symptoms ', 'Total Bilirubin(mg/dL)']
```

Checking explainer for XGB1...
<shap.explainers._tree.Tree object at 0x7f8901cf1670>

Checking shap values for XGB1...

```
[[[-4.1213792e-02 -3.3411807e-01  3.1376600e-01 ... -3.7167885e-03
    5.5054866e-02 -5.6627635e-02]
 [ 8.7244794e-02  3.1003144e-03 -1.0110403e-01 ... -8.8498723e-03
 -2.7989864e-01 -7.0158280e-02]
 [-4.6925444e-02  1.0683852e-01  3.3140031e-01 ... -7.7774939e-03
 -2.4780509e-01 -1.7681159e-04]
 ...
 [-1.6688924e-02 -1.4458889e-01  2.3780176e-01 ... -4.2031193e-03
 -2.1513927e-01 -1.9822428e-02]
 [-1.9945810e-02  1.5445979e-01  2.7486852e-01 ...  8.2303239e-03
  7.4652217e-02 -1.7326010e-02]
 [-3.5156224e-02 -2.9176781e-01  7.5178228e-02 ... -5.8821207e-03
  5.9577417e-02 -8.8170484e-02]]
```

Checking shap plots for XGB1...

Expected value for XGB: 0.40734654664993286
Summary Plot for SHAP Values in Test Set:

SHAP Bar Plot for SHAP Values Test Set:

SHAP Decision Plot for SHAP Values in Test Set:

SHAP Decision Plot for Single-Prediction in Test Set:

Checking feature importance for XGB1...

Checking for list of csv files...

```
['/Users/jessicakim/Desktop/STREAMLINE/DemoData/Output/hcc_demo/hcc-data_example_no_covariates/model_evaluation/shap_v
alues/testResults/XGB_1_shapFIVValues_Test.csv']
```

Unnamed: 0	Features	Importance
0	Haemoglobin (g/dL)	0.579048
1	Alpha-Fetoprotein (ng/mL)	0.320530
2	Oxygen Saturation (%)	0.304379
3	Performance Status*	0.301961
4	Iron	0.271850
5	Direct Bilirubin (mg/dL)	0.257519
6	International Normalised Ratio*	0.224470
7	Mean Corpuscular Volume	0.223468
8	Alkaline phosphatase (U/L)	0.198403
9	Ascites degree*	0.183599
10	Albumin (mg/dL)	0.183055
11	Ferritin (ng/mL)	0.169407
12	Symptoms	0.134469
13	Leukocytes(G/L)	0.127525
14	Platelets	0.125923
15	Aspartate transaminase (U/L)	0.117462
16	Major dimension of nodule (cm)	0.093465
17	Total Bilirubin(mg/dL)	0.061715
18	Packs of cigarets per year	0.047774
19	Alanine transaminase (U/L)	0.045326
20	Gamma glutamyl transferase (U/L)	0.035770
21	Number of Nodules	0.032060
22	Creatinine (mg/dL)	0.030768

23	23	Diabetes	0.022067
24	24	Hepatitis C Virus Antibody	0.020044
25	25	Encephalopathy degree*	0.011475
26	26	Portal Hypertension	0.007992
27	27	Arterial Hypertension	0.007300
28	28	Portal Vein Thrombosis	0.006735
29	29	Obesity	0.002671
30	30	Liver Metastasis	0.002019
31	31	Hemochromatosis	0.000000
32	32	Endemic Countries	0.000000
33	33	Cirrhosis	0.000000
34	34	Nonalcoholic Steatohepatitis	0.000000
35	35	Hepatitis B e Antigen	0.000000

Out [18]:

Unnamed: 0		Features	Importance
0	0	Haemoglobin (g/dL)	0.579048
1	1	Alpha-Fetoprotein (ng/mL)	0.320530
2	2	Oxygen Saturation (%)	0.304379
3	3	Performance Status*	0.301961
4	4	Iron	0.271850
5	5	Direct Bilirubin (mg/dL)	0.257519
6	6	International Normalised Ratio*	0.224470
7	7	Mean Corpuscular Volume	0.223468
8	8	Alkaline phosphatase (U/L)	0.198403
9	9	Ascites degree*	0.183599
10	10	Albumin (mg/dL)	0.183055
11	11	Ferritin (ng/mL)	0.169407
12	12	Symptoms	0.134469
13	13	Leukocytes(G/L)	0.127525
14	14	Platelets	0.125923
15	15	Aspartate transaminase (U/L)	0.117462
16	16	Major dimension of nodule (cm)	0.093465
17	17	Total Bilirubin(mg/dL)	0.061715
18	18	Packs of cigarets per year	0.047774
19	19	Alanine transaminase (U/L)	0.045326
20	20	Gamma glutamyl transferase (U/L)	0.035770
21	21	Number of Nodules	0.032060
22	22	Creatinine (mg/dL)	0.030768
23	23	Diabetes	0.022067
24	24	Hepatitis C Virus Antibody	0.020044
25	25	Encephalopathy degree*	0.011475
26	26	Portal Hypertension	0.007992
27	27	Arterial Hypertension	0.007300
28	28	Portal Vein Thrombosis	0.006735
29	29	Obesity	0.002671
30	30	Liver Metastasis	0.002019
31	31	Hemochromatosis	0.000000
32	32	Endemic Countries	0.000000
33	33	Cirrhosis	0.000000
34	34	Nonalcoholic Steatohepatitis	0.000000
35	35	Hepatitis B e Antigen	0.000000

XGB2 In CV2...

```
Checking if correct model is loaded...
XGBClassifier(alpha=2.925626174679448e-07, base_score=0.5, booster='gbtree',
              callbacks=None, colsample_bylevel=1, colsample_bynode=1,
              colsample_bytree=0.10192679583709846, early_stopping_rounds=None,
              enable_categorical=False, eta=0.0002501390367621765,
              eval_metric=None, gamma=4.155123430773271e-05, gpu_id=-1,
              grow_policy='depthwise', importance_type=None,
              interaction_constraints='', learning_rate=0.000250139041,
              max_bin=256, max_cat_to_onehot=4, max_delta_step=0, max_depth=2,
              max_leaves=0, min_child_weight=6.224435947630516,
              min_samples_leaf=10, min_samples_split=45, missing=nan,
              monotone_constraints=('', n_estimators=956, n_jobs=1, nthread=1, ...)
['Albumin (mg/dL)', 'Alcohol', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/mL)', 'Arterial Hypertension', 'As
cites degree*', 'Aspartate transaminase (U/L)', 'Chronic Renal Insufficiency', 'Cirrhosis', 'Creatinine (mg/dL)', 'Dia
betes', 'Direct Bilirubin (mg/dL)', 'Endemic Countries', 'Ferritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haem
oglobin (g/dL)', 'Hemochromatosis', 'Hepatitis B Surface Antigen', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibod
y', 'Human Immunodeficiency Virus', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Metastasis',
'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Nonalcoholic Steatohepatitis', 'Number of Nodules', 'Oxy
gen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Platelets', 'Portal Hypertension', 'Portal
Vein Thrombosis', 'Smoking']
['Albumin (mg/dL)', 'Alcohol', 'Alkaline phosphatase (U/L)', 'Alpha-Fetoprotein (ng/mL)', 'Arterial Hypertension', 'As
cites degree*', 'Aspartate transaminase (U/L)', 'Chronic Renal Insufficiency', 'Cirrhosis', 'Creatinine (mg/dL)', 'Dia
betes', 'Direct Bilirubin (mg/dL)', 'Endemic Countries', 'Ferritin (ng/mL)', 'Gamma glutamyl transferase (U/L)', 'Haem
oglobin (g/dL)', 'Hemochromatosis', 'Hepatitis B Surface Antigen', 'Hepatitis B e Antigen', 'Hepatitis C Virus Antibod
y', 'Human Immunodeficiency Virus', 'International Normalised Ratio*', 'Iron', 'Leukocytes(G/L)', 'Liver Metastasis',
'Major dimension of nodule (cm)', 'Mean Corpuscular Volume', 'Nonalcoholic Steatohepatitis', 'Number of Nodules', 'Oxy
gen Saturation (%)', 'Packs of cigarets per year', 'Performance Status*', 'Platelets', 'Portal Hypertension', 'Portal
Vein Thrombosis', 'Smoking']
```

Checking explainer for XGB2...
<shap.explainers._tree.Tree object at 0x7f88b0205b80>

Checking shap values for XGB2...

```
[[ 5.08398050e-03  1.09896610e-05 -1.44367898e-02 ... -2.88566196e-04
 -5.45803457e-04  5.12634986e-04]
 [ 4.83505754e-03  2.37361382e-06  1.88300461e-02 ...  5.14981046e-04
 -4.53948451e-04 -9.65057523e-04]
 [ 4.78631770e-03 -3.62699211e-05  1.88307650e-02 ... -2.88566196e-04
 -5.20780042e-04  5.26774325e-04]
 ...
 [ 4.53065475e-03  2.37361382e-06  1.87011994e-02 ... -2.88566196e-04
 -4.90862702e-04 -9.11067938e-04]
 [-1.01444125e-02  2.37361382e-06  9.76637285e-03 ... -2.88566196e-04
 -4.72357147e-04 -8.52823956e-04]
 [ 4.44364175e-03  1.09896610e-05 -2.03168858e-02 ... -2.76739825e-04
 -5.45803457e-04  5.12634986e-04]]
```

Checking shap plots for XGB2...

Expected value for XGB: -0.0014823254896327853
Summary Plot for SHAP Values in Test Set:

SHAP Bar Plot for SHAP Values Test Set:

SHAP Decision Plot for SHAP Values in Test Set:

SHAP Decision Plot for Single-Prediction in Test Set:

Checking feature importance for XGB2...

Checking for list of csv files...

```
['/Users/jessicakim/Desktop/STREAMLINE/DemoData/Output/hcc_demo/hcc-data_example_no_covariates/model_evaluation/shap_v
alues/testResults/XGB_2_shapFIValues_Test.csv']
```

Unnamed: 0	Features	Importance
0	Alkaline phosphatase (U/L)	0.017751
1	Alpha-Fetoprotein (ng/mL)	0.011837
2	Performance Status*	0.010535
3	Iron	0.010486
4	Ferritin (ng/mL)	0.010484
5	Leukocytes(G/L)	0.008156
6	Haemoglobin (g/dL)	0.008135
7	Gamma glutamyl transferase (U/L)	0.007202
8	Major dimension of nodule (cm)	0.007160
9	Albumin (mg/dL)	0.006140
10	Aspartate transaminase (U/L)	0.004504
11	Oxygen Saturation (%)	0.004186
12	Mean Corpuscular Volume	0.003277
13	Number of Nodules	0.003170
14	Platelets	0.003073
15	Diabetes	0.002804
16	Direct Bilirubin (mg/dL)	0.002523
17	Ascites degree*	0.001587
18	Liver Metastasis	0.001565
19	Creatinine (mg/dL)	0.001244
20	International Normalised Ratio*	0.001193
21	Packs of cigarets per year	0.001170
22	Arterial Hypertension	0.000834

23	23	Portal Vein Thrombosis	0.000632
24	24	Smoking	0.000632
25	25	Portal Hypertension	0.000321
26	26	Alcohol	0.000013
27	27	Endemic Countries	0.000000
28	28	Hemochromatosis	0.000000
29	29	Cirrhosis	0.000000
30	30	Chronic Renal Insufficiency	0.000000
31	31	Hepatitis B e Antigen	0.000000
32	32	Hepatitis C Virus Antibody	0.000000
33	33	Human Immunodeficiency Virus	0.000000
34	34	Nonalcoholic Steatohepatitis	0.000000
35	35	Hepatitis B Surface Antigen	0.000000

Out[18]:

Unnamed: 0		Features	Importance
0	0	Alkaline phosphatase (U/L)	0.017751
1	1	Alpha-Fetoprotein (ng/mL)	0.011837
2	2	Performance Status*	0.010535
3	3	Iron	0.010486
4	4	Ferritin (ng/mL)	0.010484
5	5	Leukocytes(G/L)	0.008156
6	6	Haemoglobin (g/dL)	0.008135
7	7	Gamma glutamyl transferase (U/L)	0.007202
8	8	Major dimension of nodule (cm)	0.007160
9	9	Albumin (mg/dL)	0.006140
10	10	Aspartate transaminase (U/L)	0.004504
11	11	Oxygen Saturation (%)	0.004186
12	12	Mean Corpuscular Volume	0.003277
13	13	Number of Nodules	0.003170
14	14	Platelets	0.003073
15	15	Diabetes	0.002804
16	16	Direct Bilirubin (mg/dL)	0.002523
17	17	Ascites degree*	0.001587
18	18	Liver Metastasis	0.001565
19	19	Creatinine (mg/dL)	0.001244
20	20	International Normalised Ratio*	0.001193
21	21	Packs of cigarets per year	0.001170
22	22	Arterial Hypertension	0.000834
23	23	Portal Vein Thrombosis	0.000632
24	24	Smoking	0.000632
25	25	Portal Hypertension	0.000321
26	26	Alcohol	0.000013
27	27	Endemic Countries	0.000000
28	28	Hemochromatosis	0.000000
29	29	Cirrhosis	0.000000
30	30	Chronic Renal Insufficiency	0.000000
31	31	Hepatitis B e Antigen	0.000000
32	32	Hepatitis C Virus Antibody	0.000000
33	33	Human Immunodeficiency Virus	0.000000
34	34	Nonalcoholic Steatohepatitis	0.000000
35	35	Hepatitis B Surface Antigen	0.000000

Run SHAP for Training Sets

Optional

- This runs on training CV Datasets that were partiioned during STREAMLINE
- User can set run_train to 'True' for comparison between training and testing sets

```
In [ ]: run_force_plots = True # parameter in run_force_plot(); set to True if user wants to display force plots for trained m
run_train = False # user can change to True to run shap values for training sets

if run_train == True:
    for each in datasets:
        print("-----")
```

```
print(each)
print("-----")
full_path = experiment_path+'/'+ each

#Make folder in experiment folder/datafolder to store all shap_values per algorithm/CV combination
if not os.path.exists(full_path+'/model_evaluation/shap_values/trainResults'):
    os.mkdir(full_path+'/model_evaluation/shap_values/trainResults')

original_headers = pd.read_csv(full_path+"/exploratory/OriginalFeatureNames.csv",sep=',').columns.values.tolist
feat_order_map = {feat:i for i, feat in enumerate(original_headers)}
print(feat_order_map)

for algorithm in algorithms: #loop through algorithms
    print(abbrev[algorithm])

    for cvCount in range(0,cv_partitions): #loop through cv's
        print('{}{} In CV{}...'.format(abbrev[algorithm], cvCount, cvCount))

        # unpickle and load model
        result_file = full_path+ '/models/pickledModels/' + abbrev[algorithm]+ "_" + str(cvCount)+".pickle"
        file = open(result_file, 'rb')
        model = pickle.load(file)
        file.close()
        print('\nChecking if correct model is loaded...\n', model)

        # Load CV datasets, paths to datasets updates with each iteration
        train_path = experiment_path + train_file_path + 'hcc-data_example_CV_' + str(cvCount) + '_Train.csv'
        test_path = experiment_path + test_file_path + 'hcc-data_example_CV_' + str(cvCount) + '_Test.csv'
        trainX, trainY,testX, testY, train_feat, test_feat = dataPrep(train_path,instance_label,class_label, t

        # shap computation and plots
        explainer = get_explainer(model, abbrev[algorithm], trainX)
        print('\nChecking explainer for {}{}...\n{}'.format(abbrev[algorithm], cvCount, explainer)) # print e

        print('\nChecking shap values for {}{}...\n'.format(abbrev[algorithm], cvCount))
        shap_values = compute_shapValues(model, abbrev[algorithm], explainer, trainX)

        print('\nChecking shap plots for {}{}...\n'.format(abbrev[algorithm], cvCount))
        shap_summary(abbrev[algorithm], train_feat, shap_values, explainer, trainX)

        #save SHAP FI results
        print('\nChecking feature importance for {}{}...\n'.format(abbrev[algorithm], cvCount))
        shap_fi_df = shap_feature_ranking(abbrev[algorithm], shap_values, trainX, train_feat) # can either cho

        filepath = full_path+"/model_evaluation/shap_values/trainResults/"+ abbrev[algorithm] + '_' + str(cvCo
        shap_fi_df.to_csv(filepath, header=True, index=True)

        # only runs force plots if run = True
        if run_force_plots == True:
            if abbrev[algorithm] in ['NB']:

                print('\nForce Plot for {}{} SHAP Values in Train Set: \n'.format(abbrev[algorithm], cvCount))
                shap.force_plot(shap_values, trainX, feature_names=train_feat)

                print('\nSingle-Prediction Force Plot for {}{} SHAP Values in Train Set: \n'.format(abbrev[alg
                shap.force_plot(shap_values[42], trainX.iloc[42], feature_names=train_feat, show=False)
                plt.savefig(full_path+'/model_evaluation/'+abbrev[algorithm]+"_shapFP.png",dpi=300) FIXME
                break

            elif abbrev[algorithm] in ['LR', 'XGB', 'LGB', 'CBG']: #need to test out LGB and CBG for this

                print('\nForce Plot for {}{} SHAP Values in Whole Train Set: \n'.format(abbrev[algorithm], cv
                shap.force_plot(explainer.expected_value, shap_values, trainX, feature_names=train_feat)

                print('\nSingle-Prediction Force Plot for {}{} SHAP Values in Train Set: \n'.format(abbrev[alg
                shap.force_plot(explainer.expected_value, shap_values[42], trainX.iloc[42], feature_names=trai
                break

            else:

                print('\nForce Plot for {}{} SHAP Values from Class 0 in Train Set: \n'.format(abbrev[algorith
                shap.force_plot(explainer.expected_value[0], shap_values[0], feature_names=train_feat)

                print('\nForce Plot for {}{} SHAP Values from Class 1 in Train Set: \n'.format(abbrev[algorith
                shap.force_plot(explainer.expected_value[1], shap_values[1], feature_names=train_feat)
                break
```

In []:

In []: