

```
In [7]: import pandas as pd
import numpy as np
from lifelines import CoxPHFitter
import matplotlib.pyplot as plt

# Load the data from the Excel file
data = pd.read_excel('data1.xlsx')

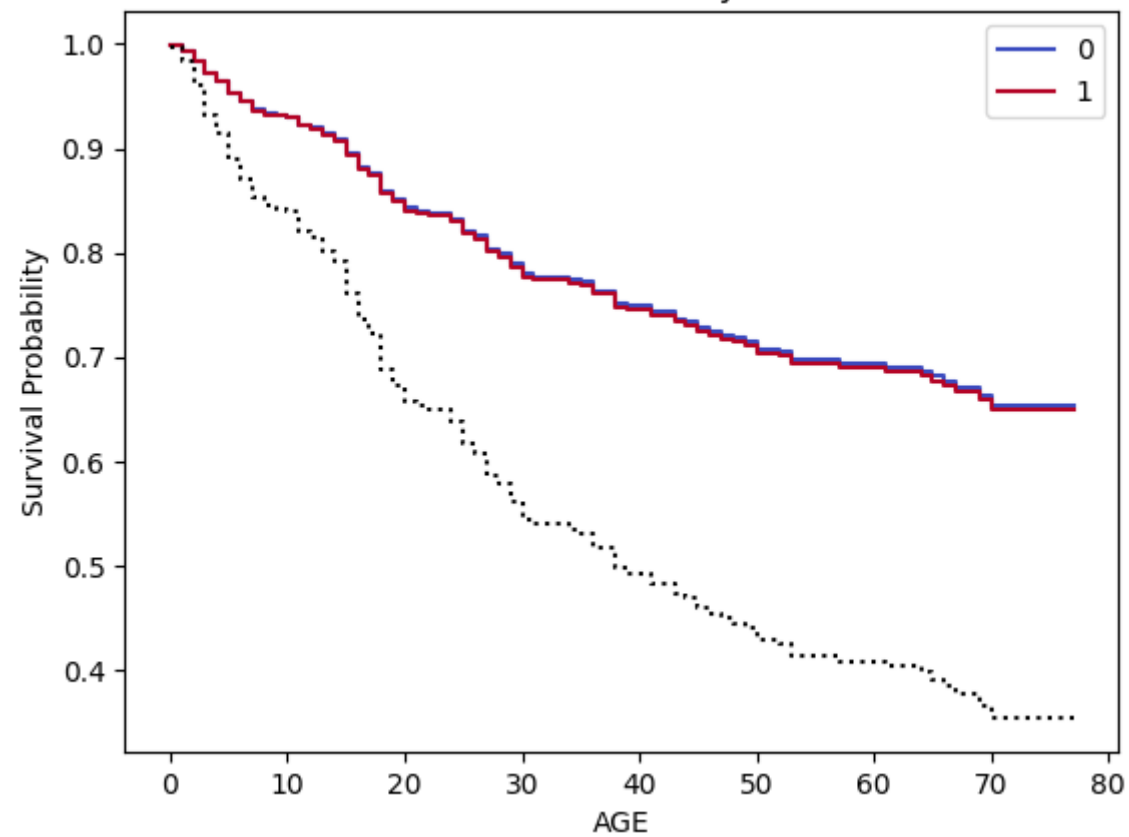
# Create a new instance of the CoxPHFitter class
cph = CoxPHFitter()

# Fit the Cox Proportional Hazard model to the data
cph.fit(data, duration_col='Months', event_col='DEATH')
```

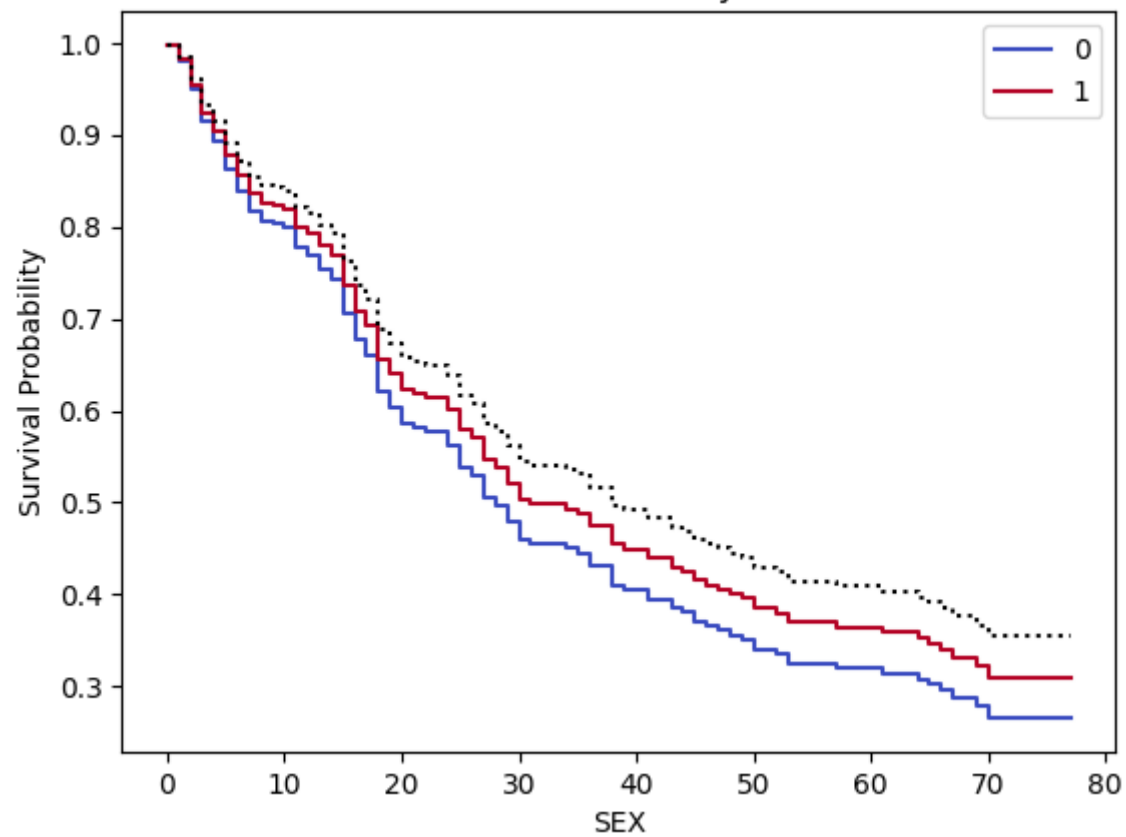
Out[7]: <lifelines.CoxPHFitter: fitted with 343 total observations, 167 right-censored observations>

```
In [2]: covariates_to_plot = ['AGE', 'SEX', 'CompositeStage', 'LNInvolment', 'Comorbidity', 'FamilyHistoryOfCancer']
for covariate in covariates_to_plot:
    cph.plot_partial_effects_on_outcome(covariates=covariate, values=[0, 1], cmap='coolwarm')
    plt.xlabel(covariate)
    plt.ylabel('Survival Probability')
    plt.title('Survival Curves by ' + covariate)
    plt.legend(['0', '1'])
    plt.show()
```

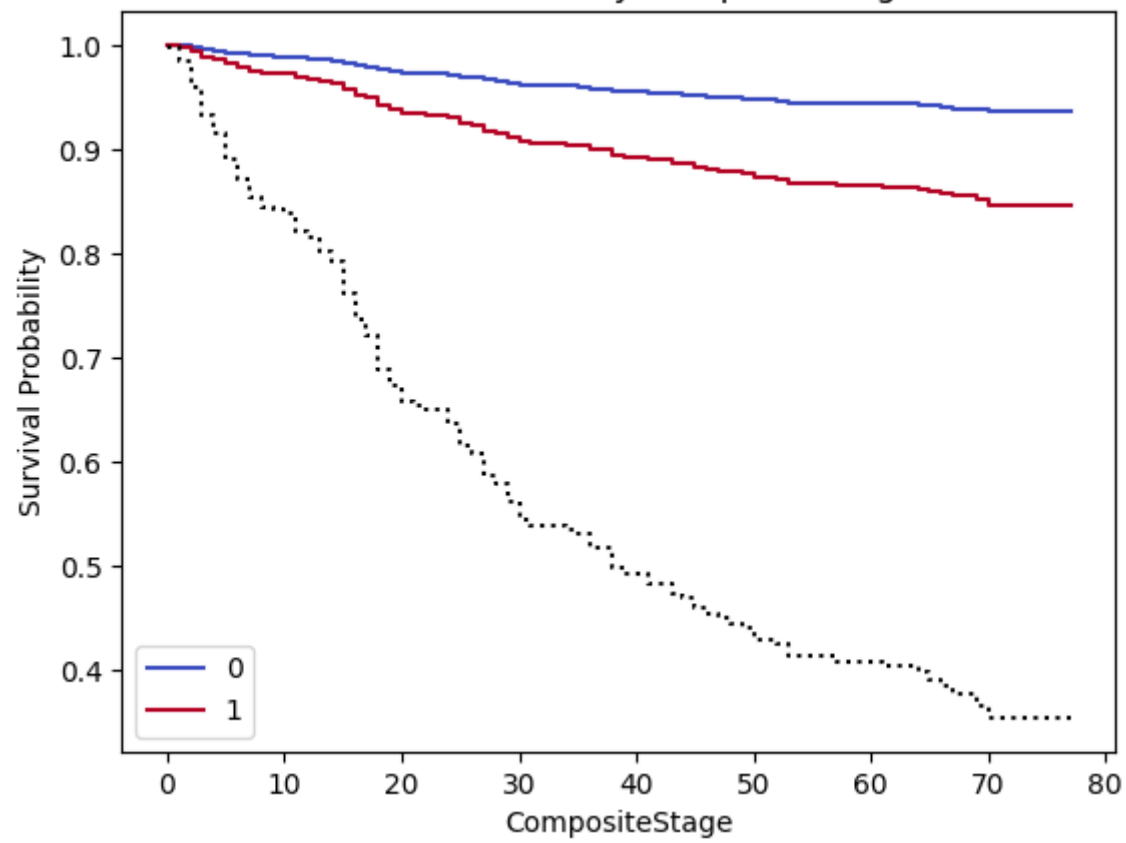
Survival Curves by AGE



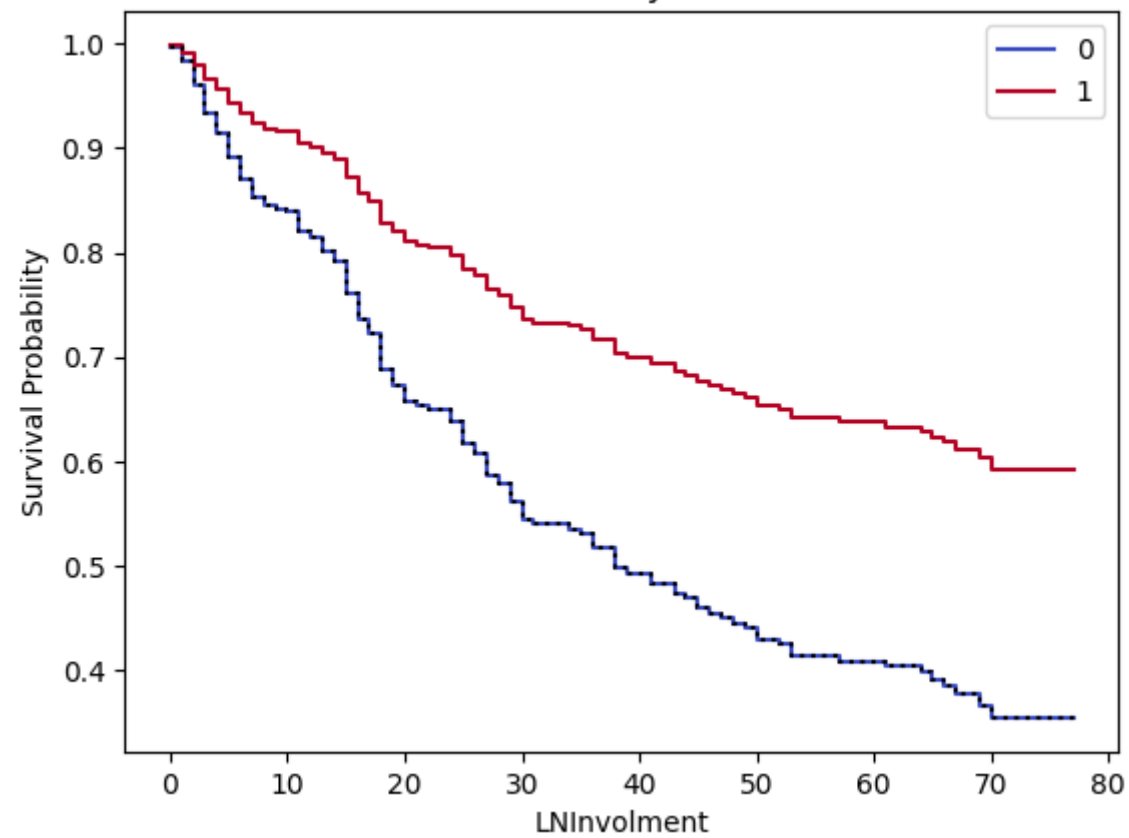
Survival Curves by SEX



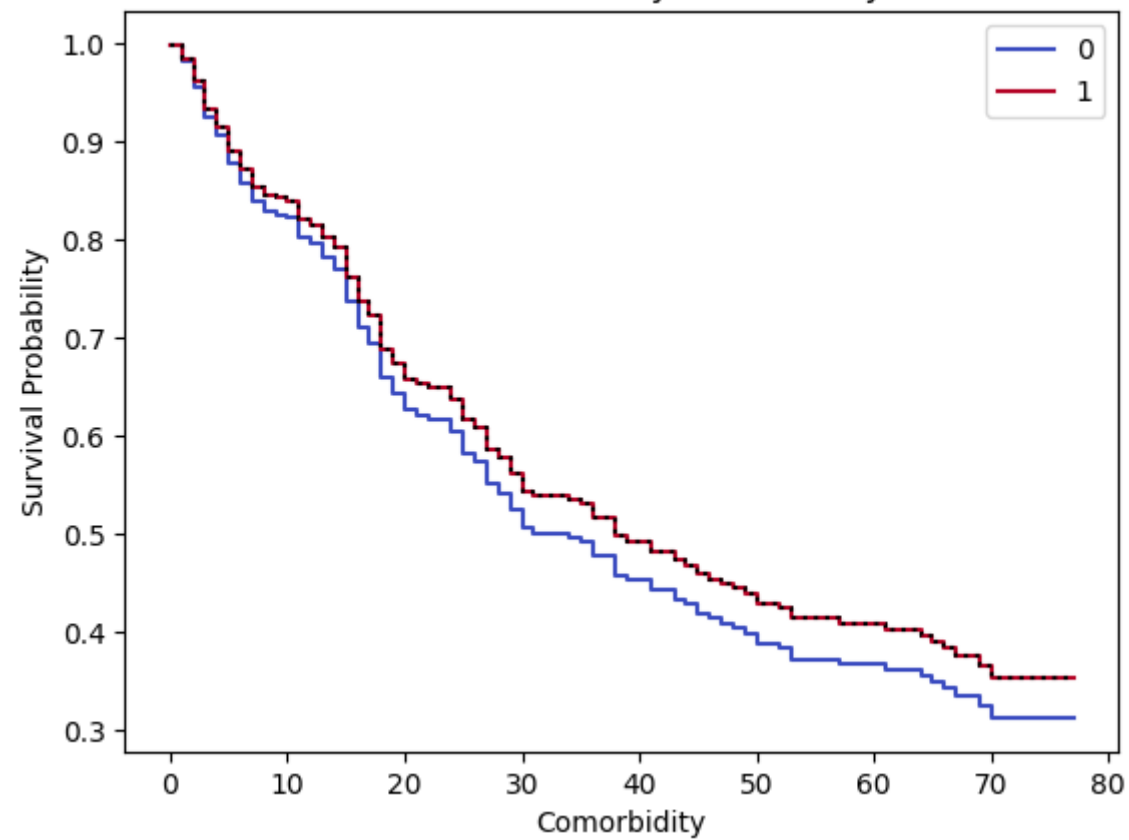
Survival Curves by CompositeStage

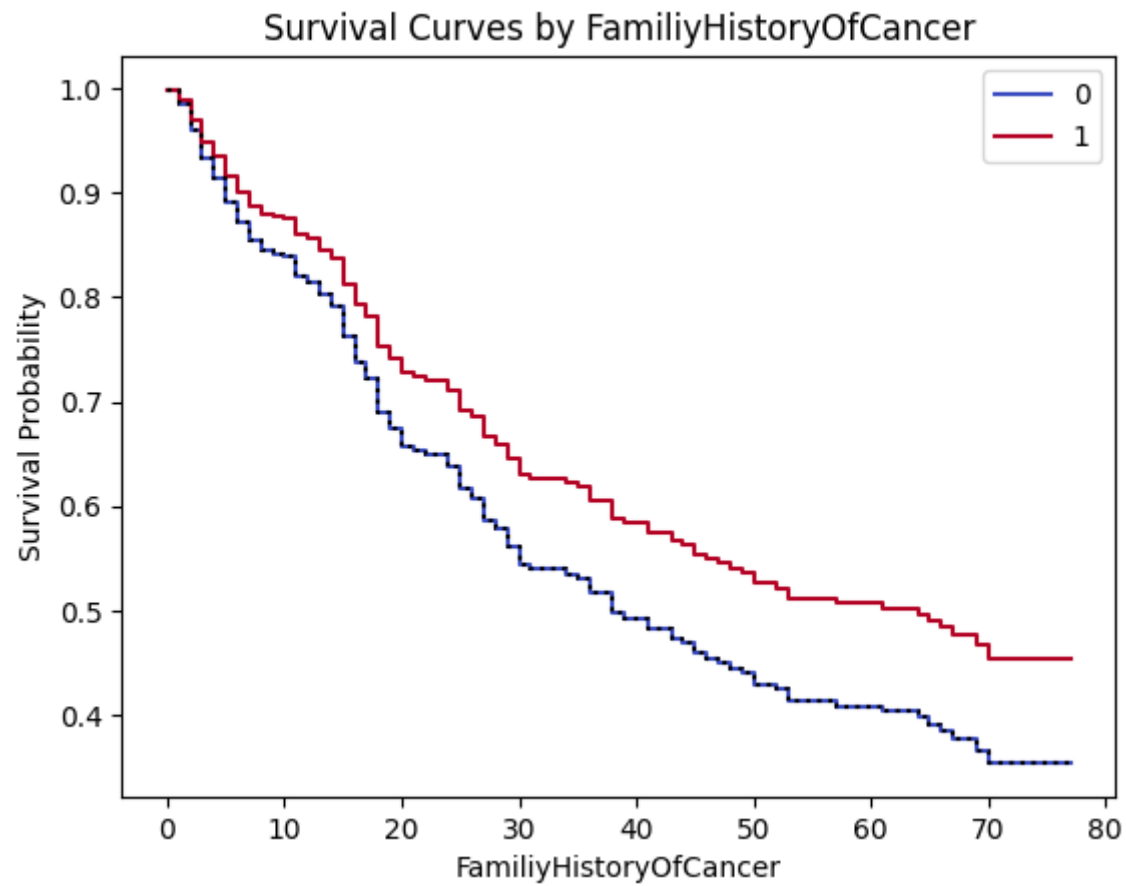


Survival Curves by LNInvolment

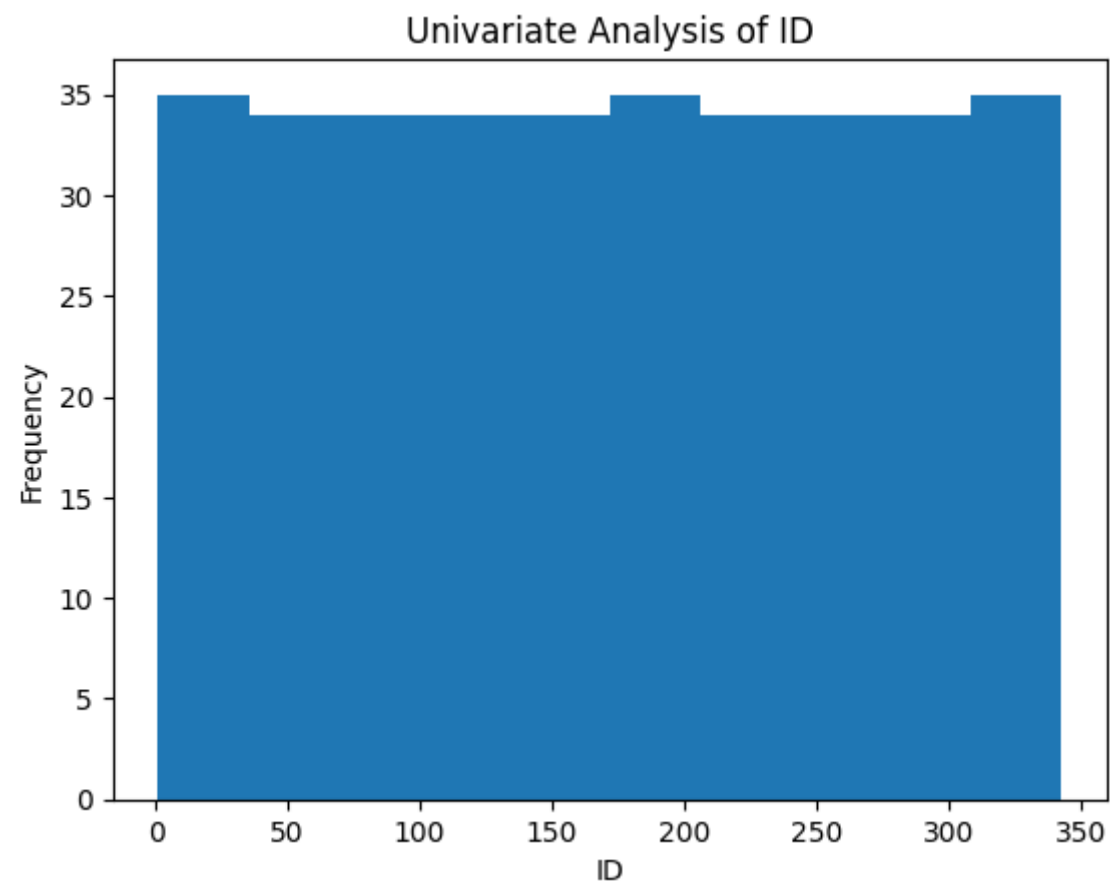


Survival Curves by Comorbidity



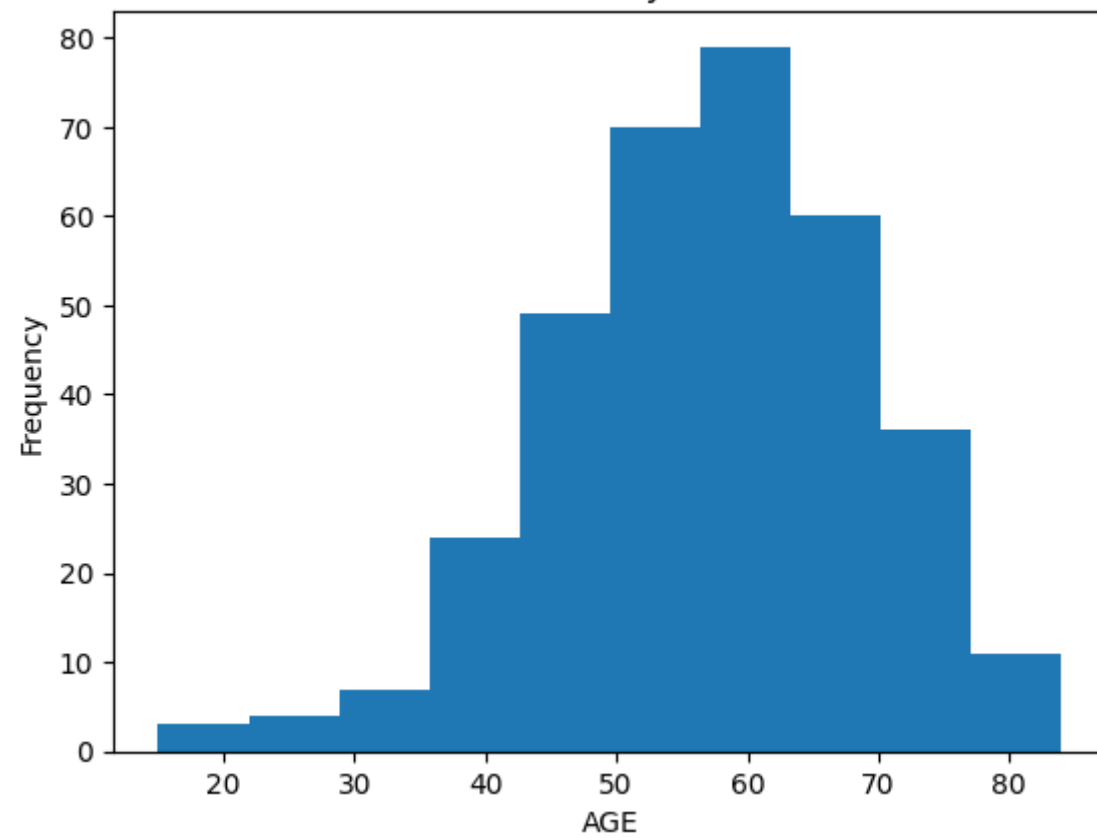


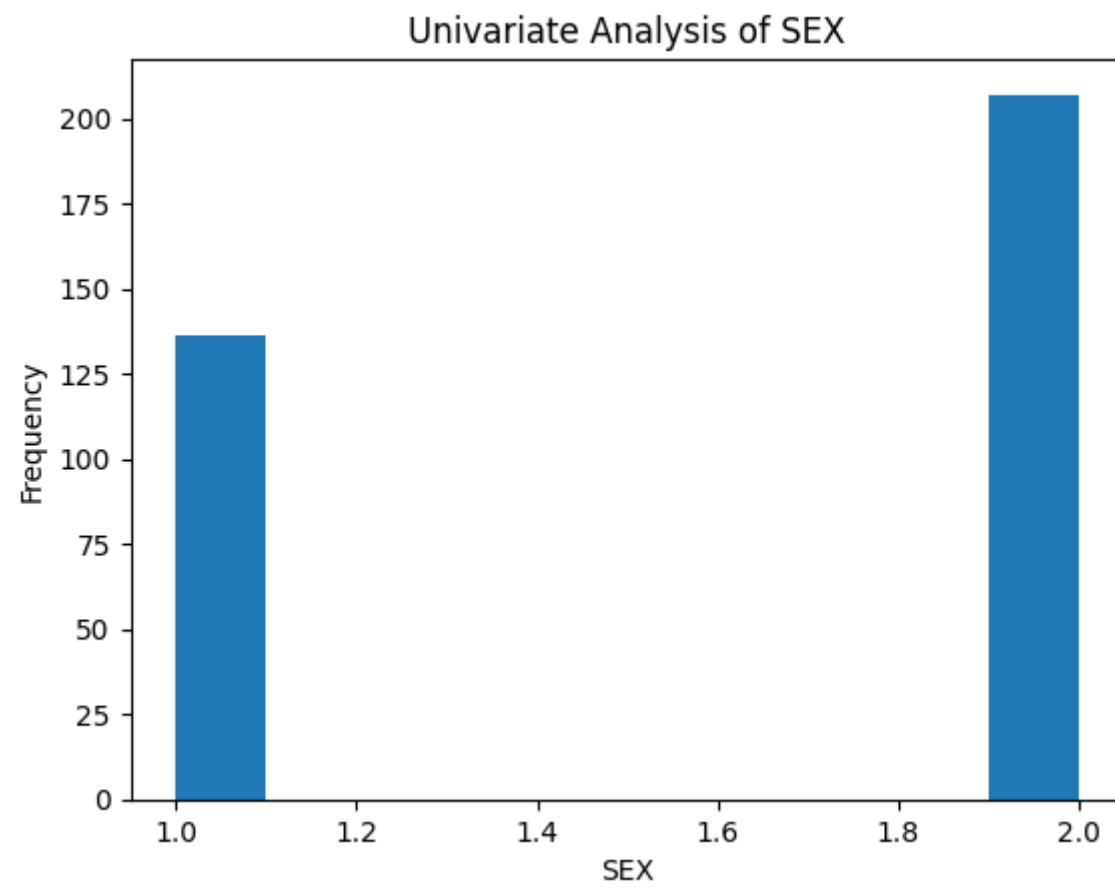
```
In [3]: for column in data.columns:
        if column not in ['Months', 'DEATH']:
            # Plot the variable
            plt.figure()
            plt.hist(data[column])
            plt.xlabel(column)
            plt.ylabel('Frequency')
            plt.title(f'Univariate Analysis of {column}')
            plt.show()
```



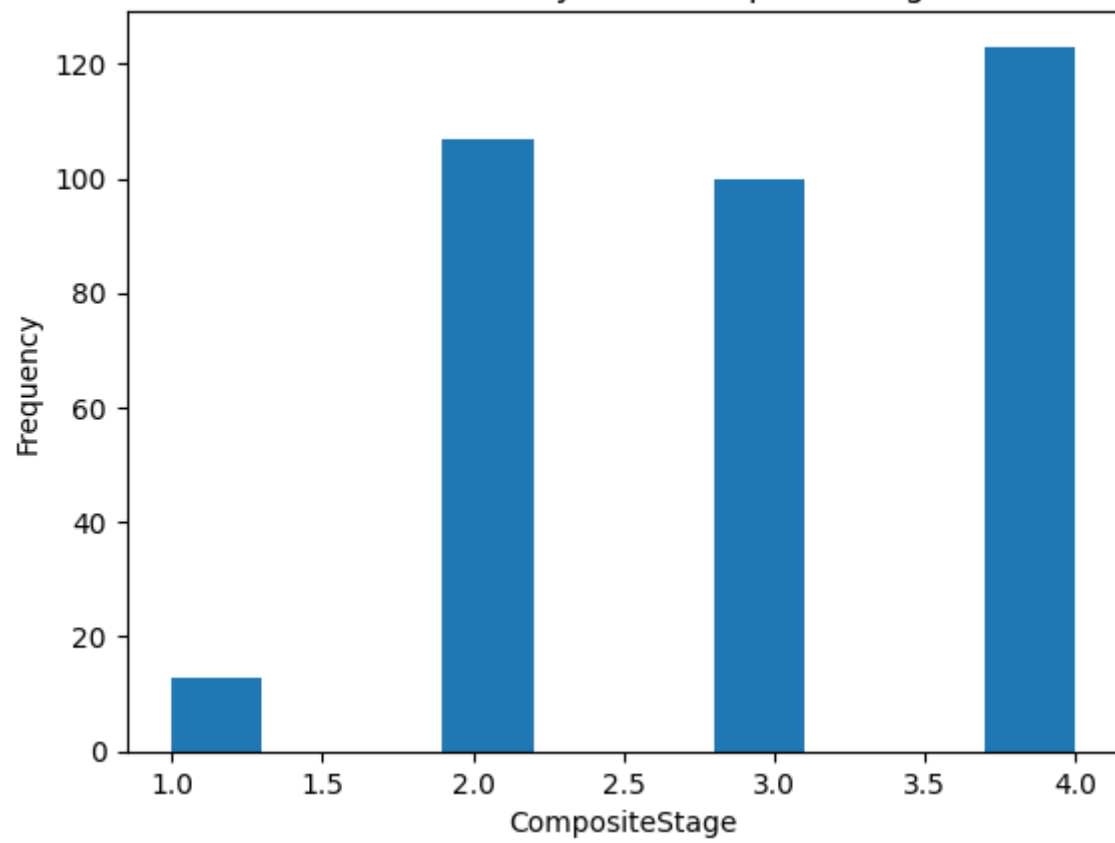


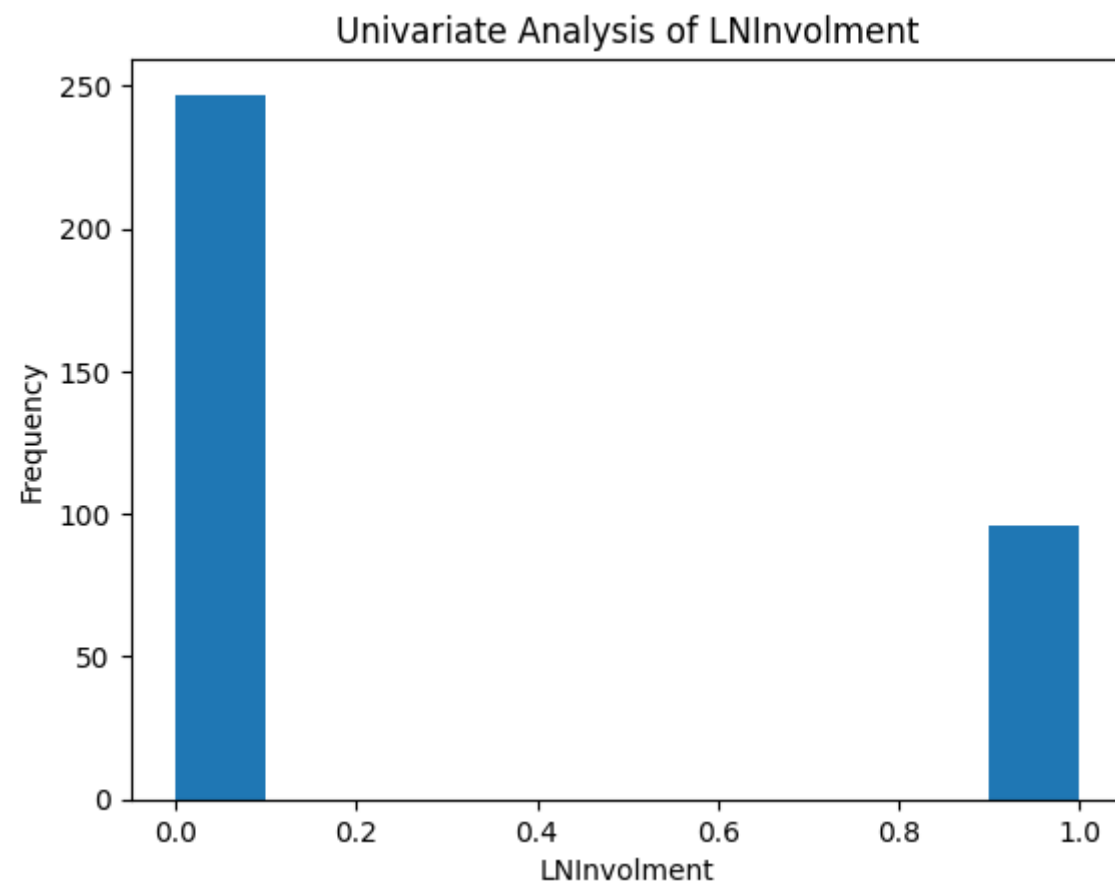
Univariate Analysis of AGE



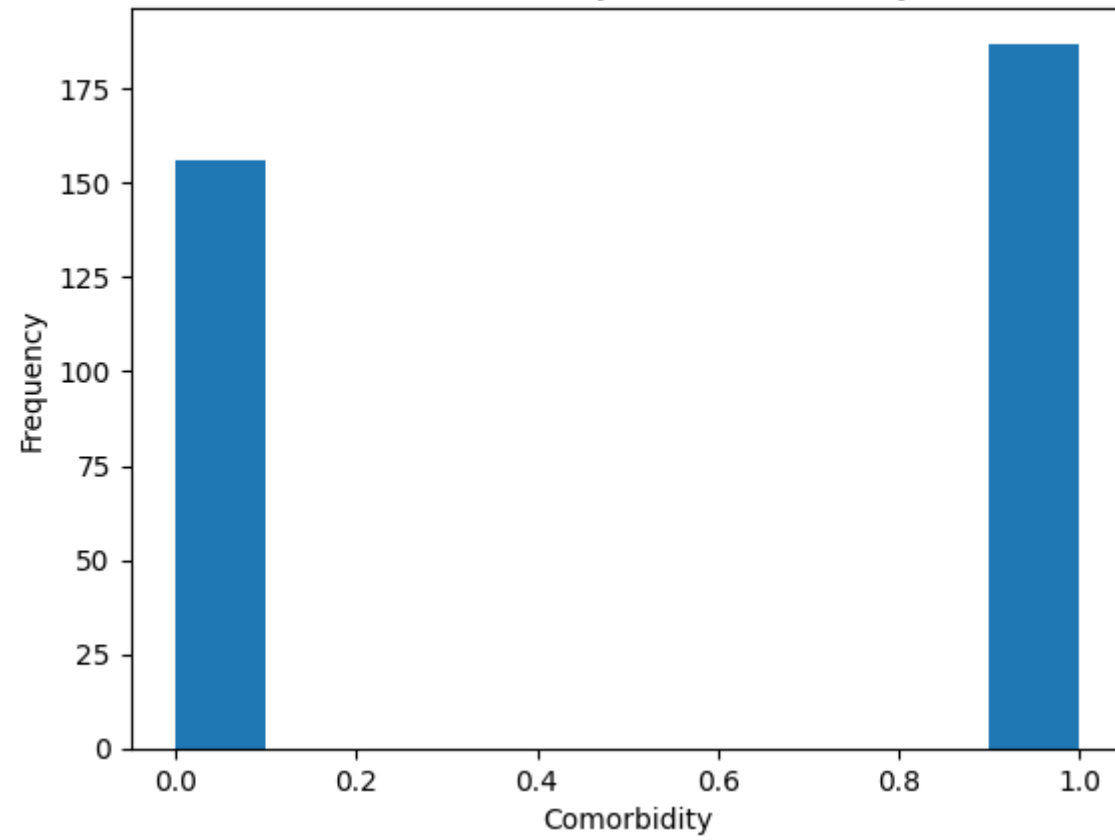


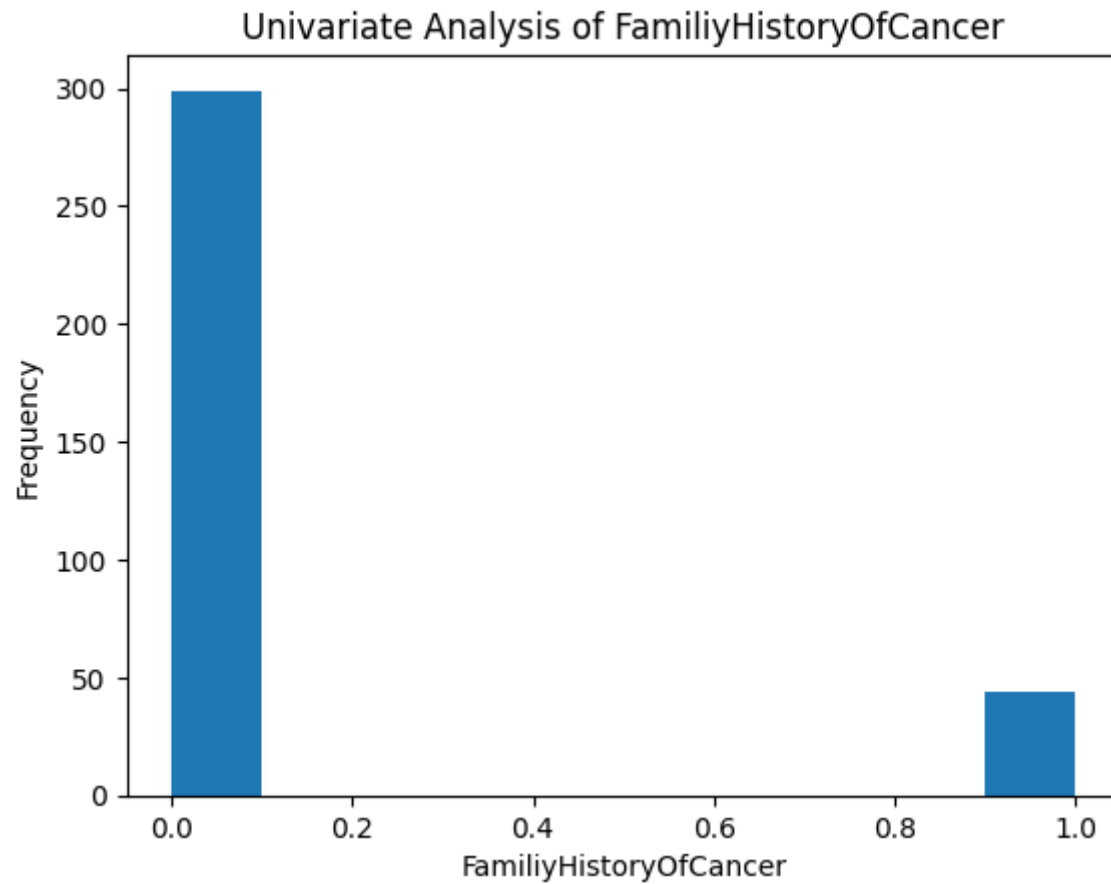
Univariate Analysis of CompositeStage





Univariate Analysis of Comorbidity





```
In [5]: n = len(data)
llf = cph.log_likelihood_
k = cph.params_.shape[0]
aic = -2 * llf + 2 * k
bic = -2 * llf + k * np.log(n)
print("AIC of the above given data:", aic)
print("BIC of the above given data:", bic)
```

AIC of the above given data: 1769.942369337933  
BIC of the above given data: 1796.8064824680946

```
In [12]: import pandas as pd
import numpy as np
```

```

from lifelines import CoxPHFitter
import matplotlib.pyplot as plt

# Load the data from the Excel file
data = pd.read_excel('data1.xlsx')

# Create a new instance of the CoxPHFitter class
cph = CoxPHFitter()

# Fit the Cox Proportional Hazard model to the data
cph.fit(data, duration_col='Months', event_col='DEATH')

covariates_to_plot = ['AGE', 'SEX', 'CompositeStage', 'LNInvolment', 'Comorbidity', 'FamilyHistoryOfCancer']
significant_covariates = []

for covariate in covariates_to_plot:
    cph.plot_partial_effects_on_outcome(covariates=covariate, values=[0, 1], cmap='coolwarm')
    plt.xlabel(covariate)
    plt.ylabel('Survival Probability')
    plt.title('Survival Curves by ' + covariate)
    plt.legend(['0', '1'])
    plt.show()

    # Prompt user to input whether the covariate is significant or not
    is_significant = input("Is " + covariate + " significant? (y/n): ")

    if is_significant.lower() == 'y':
        significant_covariates.append(covariate)

# Perform multivariate analysis
multivariate_data = data[significant_covariates + ['Months', 'DEATH']]
cph_multivariate = CoxPHFitter()
cph_multivariate.fit(multivariate_data, duration_col='Months', event_col='DEATH')

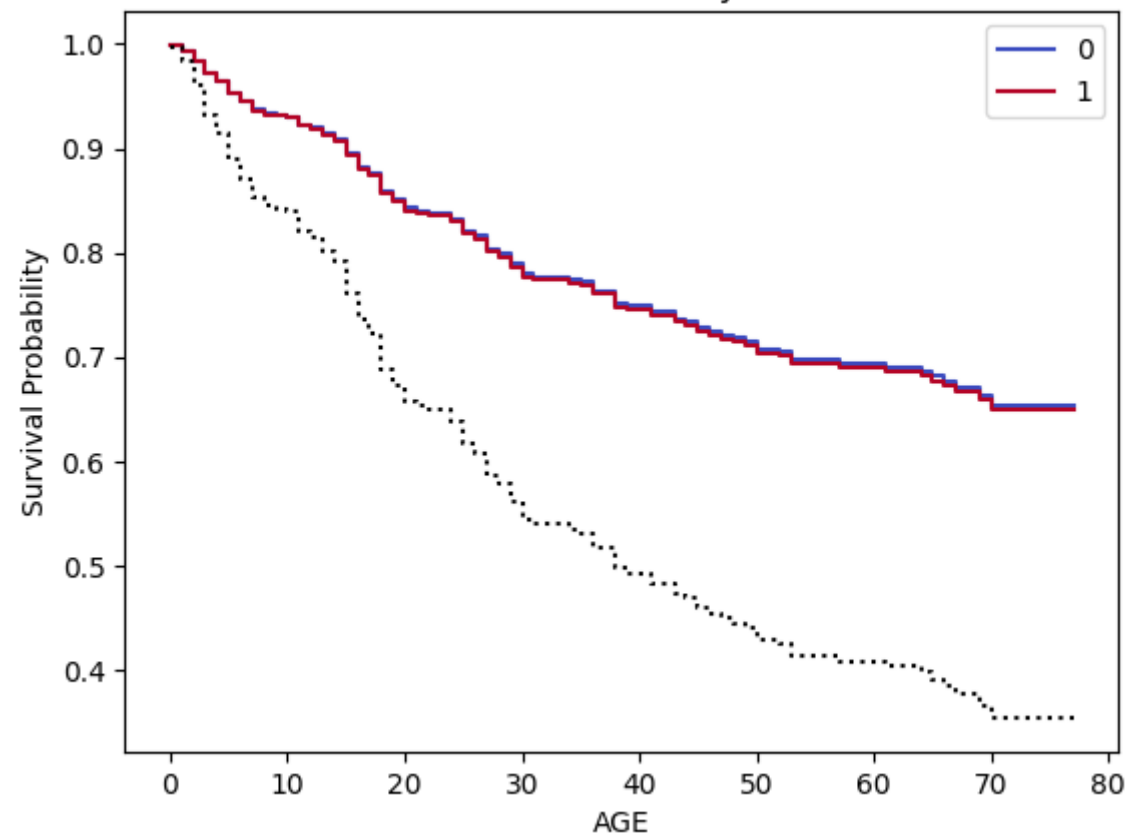
n = len(data)
llf_univariate = cph.log_likelihood_
k_univariate = cph.params_.shape[0]
aic_univariate = -2 * llf_univariate + 2 * k_univariate
bic_univariate = -2 * llf_univariate + k_univariate * np.log(n)

```

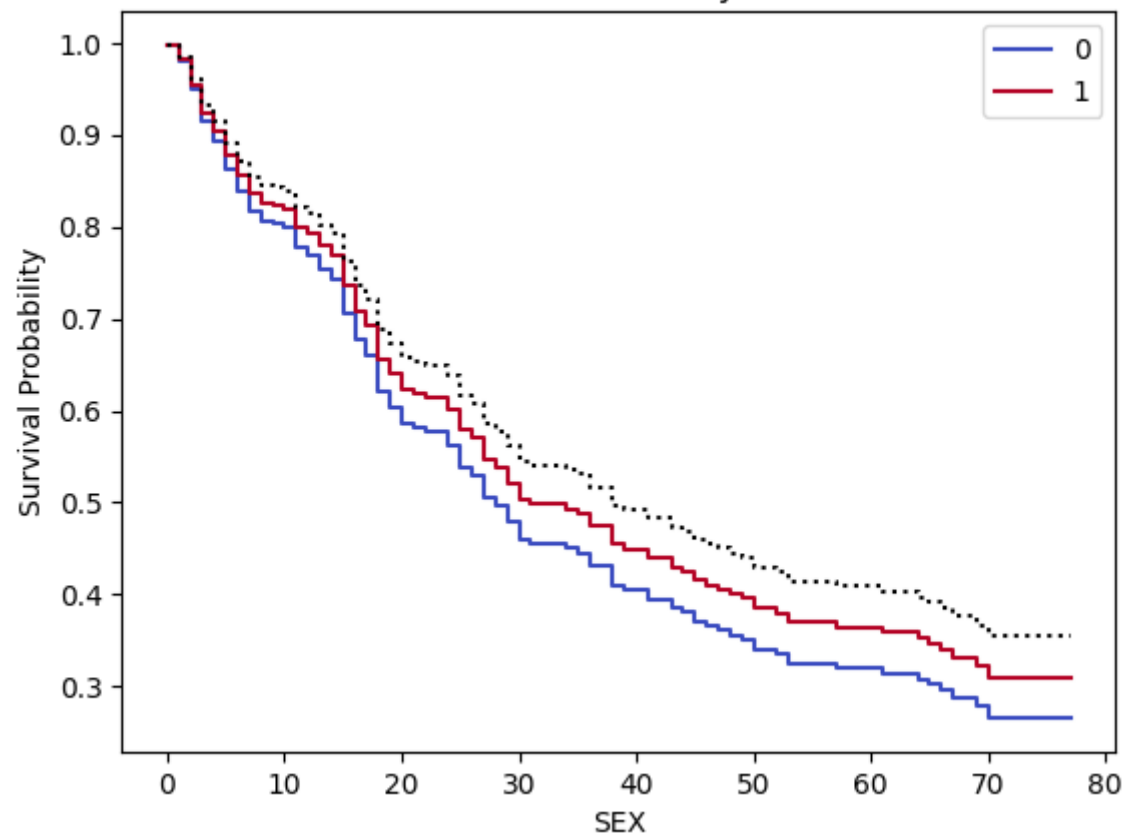
```
llf_multivariate = cph_multivariate.log_likelihood_  
k_multivariate = cph_multivariate.params_.shape[0]  
aic_multivariate = -2 * llf_multivariate + 2 * k_multivariate  
bic_multivariate = -2 * llf_multivariate + k_multivariate * np.log(n)  
  
print("Univariate Analysis:")  
print("Significant Covariates:", significant_covariates)  
print("AIC of the univariate model:", aic_univariate)  
print("BIC of the univariate model:", bic_univariate)  
  
print("\nMultivariate Analysis:")  
print(cph_multivariate.summary)  
print("AIC of the multivariate model:", aic_multivariate)  
print("BIC of the multivariate model:", bic_multivariate)
```



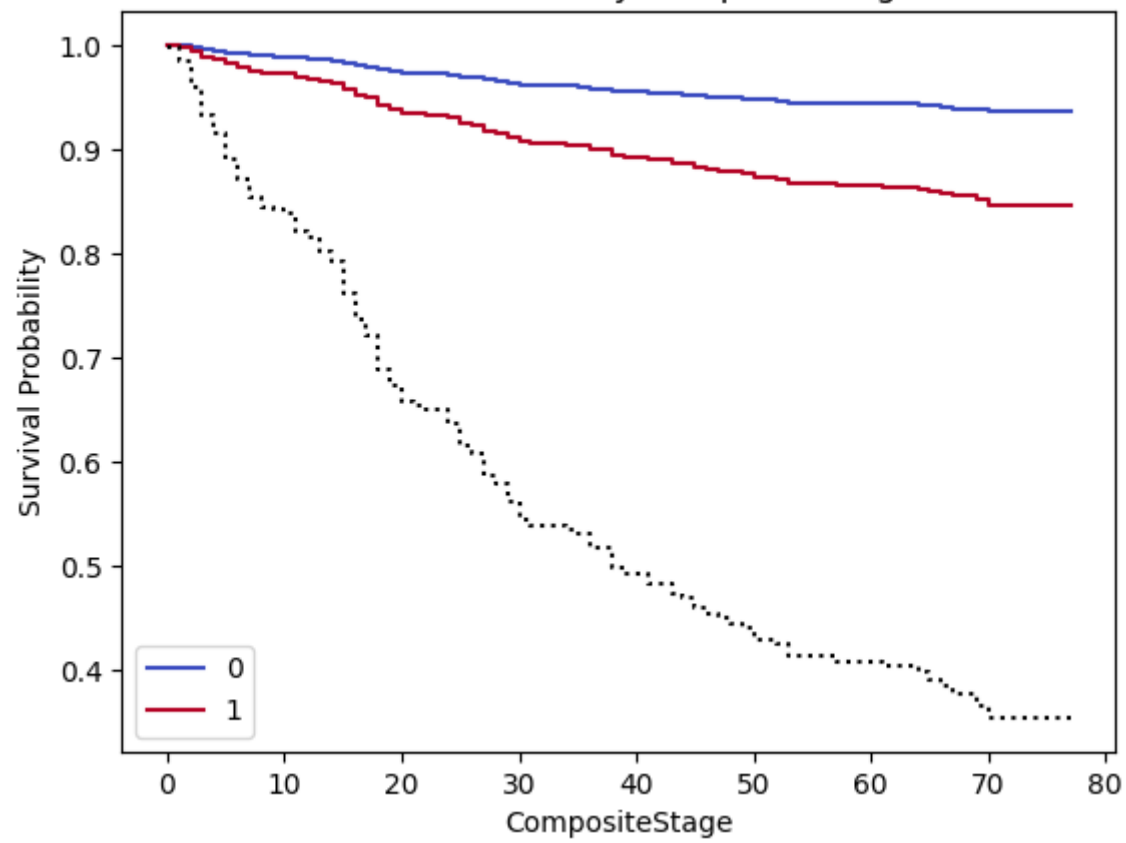
Survival Curves by AGE



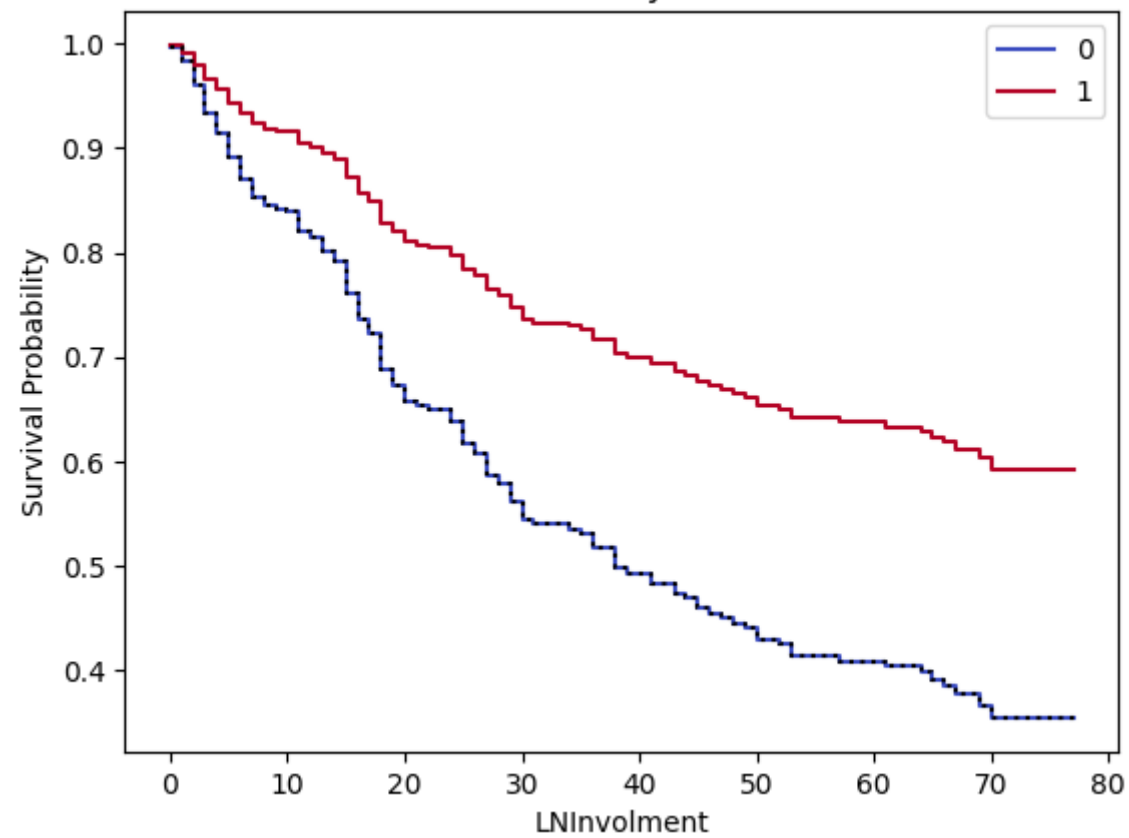
Survival Curves by SEX



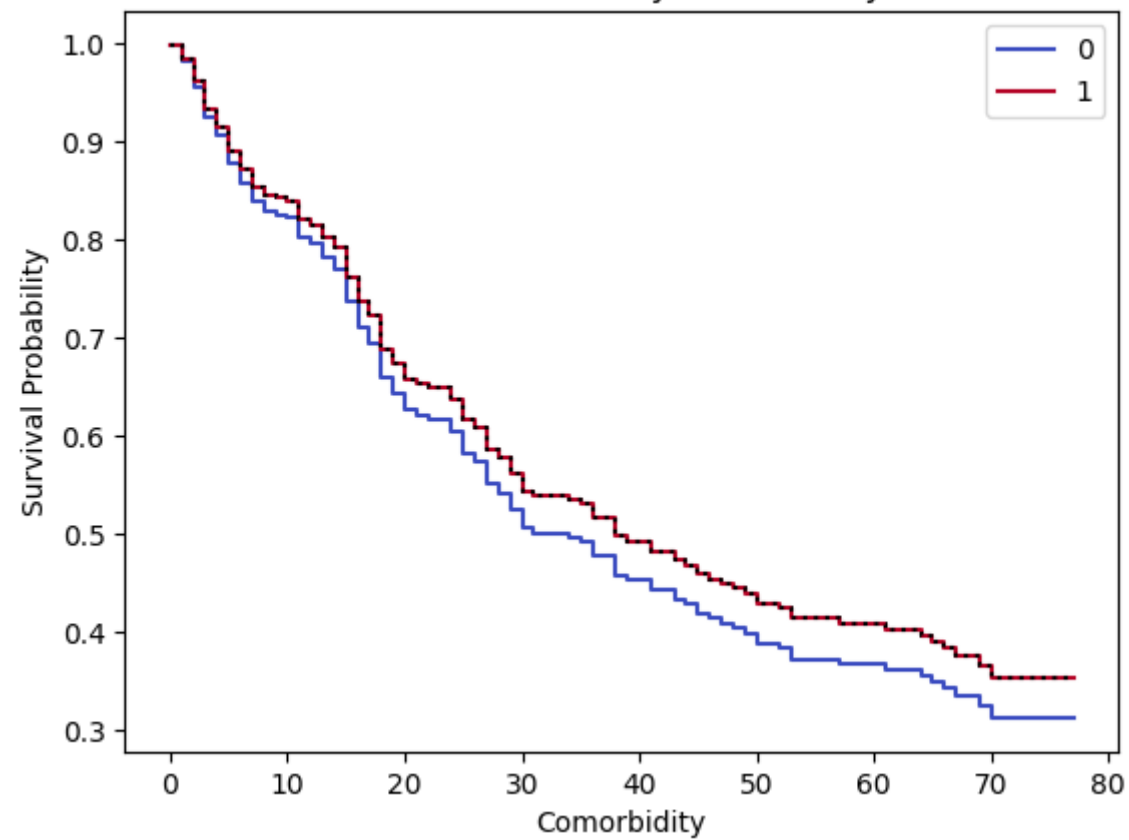
Survival Curves by CompositeStage



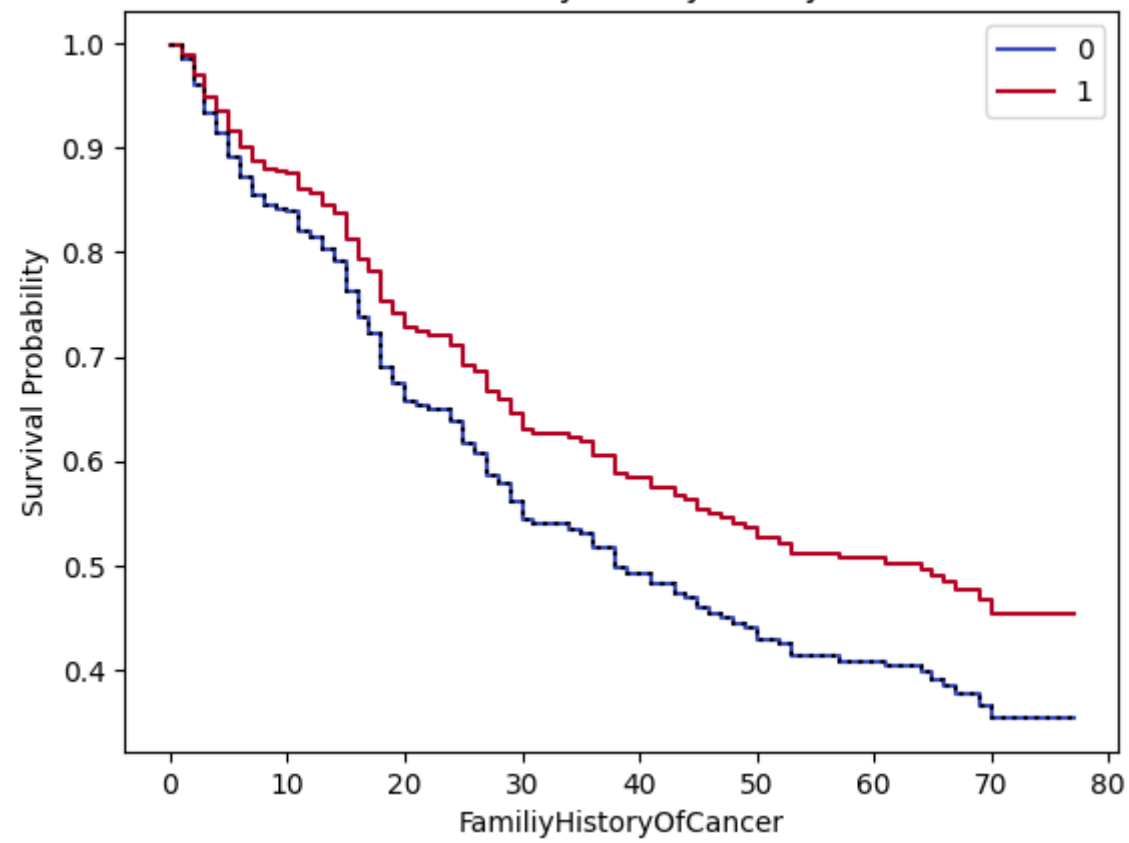
Survival Curves by LNInvolment



Survival Curves by Comorbidity



Survival Curves by FamilyHistoryOfCancer



Univariate Analysis:

Significant Covariates: ['AGE']

AIC of the univariate model: 1769.942369337933

BIC of the univariate model: 1796.8064824680946

Multivariate Analysis:

	coef	exp(coef)	se(coef)	coef lower 95%	coef upper 95%	\
covariate						
AGE	0.009362	1.009406	0.006565	-0.003506	0.02223	

	exp(coef)	lower 95%	exp(coef)	upper 95%	cmp to	z	\
covariate							
AGE		0.9965		1.022479	0.0	1.42599	

	p	-log2(p)
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covariate
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AGE	0.153871	2.700206
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AIC of the multivariate model: 1887.3267837410287

BIC of the multivariate model: 1891.1645141881947

In [ ]: