Survival Analysis

Summary Statistics

Kaplan Meier Curve

Cox Model

Weibull Regression Mode

Fit a Customized Model

Survival Probability Estimation

```
Outcome
  Dead to Breast Cancer
  (DOBC)
Variable 1
  Mutation Status
Variable 2
  NULL
Variable 3
  NULL
Variable 4
  NULL
Variable 5
 NULL
Variable 6
  NULL
Variable 7
  NULL
```

The default cox model has default outcome as Dead to Breast Cancer and default variables: Mutation Status, Age, Tumor Stage, and Node Invovlement. If wish to customize it, you can build the new model in the first column. Otherwise, the survival probability calculation will use the default.

```
Call:
coxph(formula = Surv(SURVIVAL, NDOBC) ~ ANY_MUT + NUNDER_50 +
   NT_STAGE + NNODE_ANY, data = Data)
 n= 273, number of events= 35
  (32 observations deleted due to missingness)
           coef exp(coef) se(coef)
                                    z Pr(>|z|)
ANY_MUT
         0.5685
                1.7656 0.4895 1.161 0.2455
NUNDER_50 0.2495
                  1.2833
                          0.3679 0.678
                                        0.4977
NT_STAGE 0.8959 2.4496 0.3502 2.558 0.0105 *
NNODE ANY 0.6988 2.0112 0.3573 1.955 0.0505 .
___
Signif. codes: 0 (***, 0.001 (**, 0.05 (., 0.1 (, 1
         exp(coef) exp(-coef) lower .95 upper .95
ANY_MUT
            1.766
                      0.5664
                               0.6765
                                         4.608
            1.283
                      0.7792
                               0.6240
                                         2.639
NUNDER_50
NT_STAGE
            2.450
                      0.4082
                                         4.866
                               1.2330
NNODE_ANY
                      0.4972
                                         4.052
            2.011
                               0.9984
Concordance= 0.705 (se = 0.044)
Likelihood ratio test= 16.57 on 4 df,
                                     p=0.002
Wald test
                   = 18.45 on 4 df,
                                    p=0.001
Score (logrank) test = 20.15 on 4 df,
                                     p=5e-04
```

The AIC of this model is 362.16 and the BIC of this model is 376.6 The p-value of the PH assumption is 0.28 and thus the PH assumption of this model is met.

