

The PHREG Procedure

Model Information

| | |
|--------------------|-------------------------|
| Data Set | WORK.IGAN_ANALYSIS_RANK |
| Dependent Variable | follow_up_period |
| Censoring Variable | primary_outcome |
| Censoring Value(s) | 0 |
| Ties Handling | BRESLOW |

| | |
|-----------------------------|------|
| Number of Observations Read | 1596 |
| Number of Observations Used | 1596 |

Class Level Information

| Class | Value | Design Variables |
|--------|-------|------------------|
| gender | 0 | 0 |
| | 1 | 1 |
| UA_Q4 | 0 | 0 0 0 |
| | 1 | 1 0 0 |
| | 2 | 0 1 0 |
| | 3 | 0 0 1 |

Summary of the Number of Event and Censored Values

| Total | Event | Censored | Percent Censored |
|-------|-------|----------|------------------|
| 1596 | 317 | 1279 | 80.14 |

Convergence Status

Convergence criterion (GCONV=1E-8) satisfied.

Model Fit Statistics

| Criterion | Without Covariates | With Covariates |
|-----------|--------------------|-----------------|
| -2 LOG L | 4175.145 | 4125.228 |
| AIC | 4175.145 | 4141.228 |
| SBC | 4175.145 | 4171.300 |

The PHREG Procedure

Testing Global Null Hypothesis: BETA=0

| Test | Chi-Square | DF | Pr > ChiSq |
|------------------|------------|----|------------|
| Likelihood Ratio | 49.9162 | 8 | <.0001 |
| Score | 52.9753 | 8 | <.0001 |
| Wald | 51.6468 | 8 | <.0001 |

Type 3 Tests

| Effect | DF | Wald | |
|-----------------|----|------------|------------|
| | | Chi-Square | Pr > ChiSq |
| UA_Q4 | 3 | 7.6939 | 0.0528 |
| gender | 1 | 2.7634 | 0.0964 |
| age | 1 | 0.3263 | 0.5679 |
| Proteinuria24h | 1 | 0.3625 | 0.5471 |
| GFR_base_CKDEPI | 1 | 17.1770 | <.0001 |
| MBP | 1 | 0.3518 | 0.5531 |

Analysis of Maximum Likelihood Estimates

| Parameter | DF | Parameter Estimate | Standard Error | Chi-Square | Pr > ChiSq | Hazard Ratio | 95% Hazard Ratio Confidence Limits |
|-----------------|----|--------------------|----------------|------------|------------|--------------|------------------------------------|
| UA_Q4 | 1 | 0.19991 | 0.18761 | 1.1354 | 0.2866 | 1.221 | 0.846 1.764 |
| UA_Q4 | 2 | 0.30813 | 0.19499 | 2.4970 | 0.1141 | 1.361 | 0.929 1.994 |
| UA_Q4 | 3 | 0.54927 | 0.20629 | 7.0897 | 0.0078 | 1.732 | 1.156 2.595 |
| gender | 1 | -0.21679 | 0.13041 | 2.7634 | 0.0964 | 0.805 | 0.624 1.040 |
| age | 1 | -0.00345 | 0.00605 | 0.3263 | 0.5679 | 0.997 | 0.985 1.008 |
| Proteinuria24h | 1 | 0.00439 | 0.00730 | 0.3625 | 0.5471 | 1.004 | 0.990 1.019 |
| GFR_base_CKDEPI | 1 | -0.01024 | 0.00247 | 17.1770 | <.0001 | 0.990 | 0.985 0.995 |
| MBP | 1 | -0.00278 | 0.00468 | 0.3518 | 0.5531 | 0.997 | 0.988 1.006 |

Analysis of Maximum Likelihood Estimates

| Parameter | Label |
|-----------------|--------------------------|
| UA_Q4 | 1 Rank for Variable UA 1 |
| UA_Q4 | 2 Rank for Variable UA 2 |
| UA_Q4 | 3 Rank for Variable UA 3 |
| gender | 1 gender 1 |
| age | |
| Proteinuria24h | |
| GFR_base_CKDEPI | |
| MBP | |