

## 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	1482

## Class Level Information

Class	Value	Design Variables		
gender	0	0		
	1	1		
M	0	0		
	1	1		
E	0	0		
	1	1		
S	0	0		
	1	1		
T	0	0	0	
	1	1	0	
	2	0	1	
C	0	0		
	1	1		
ACEI_ARB	0	0		
	1	1		
steroid	0	0		
	1	1		
UA_Q4	0	0	0	0
	1	1	0	0
	2	0	1	0
	3	0	0	1

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## Summary of the Number of Event and Censored Values

Total	Event	Censored	Percent Censored
1482	283	1199	80.90

## Convergence Status

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

## Model Fit Statistics

Criterion	Without Covariates	With Covariates
-2 LOG L	3674.676	3463.191
AIC	3674.676	3495.191
SBC	3674.676	3553.518

## Testing Global Null Hypothesis: BETA=0

Test	Chi-Square	DF	Pr > ChiSq
Likelihood Ratio	211.4846	16	<.0001
Score	249.3689	16	<.0001
Wald	209.4713	16	<.0001

## Type 3 Tests

Effect	DF	Wald Chi-Square	Pr > ChiSq
UA_Q4	3	14.5040	0.0023
gender	1	3.5967	0.0579
age	1	1.0083	0.3153
Proteinuria24h	1	0.0498	0.8234
GFR_base_CKDEPI	1	6.8209	0.0090
MBP	1	0.2655	0.6064
M	1	74.0937	<.0001
E	1	4.2953	0.0382
S	1	6.4055	0.0114
T	2	7.2779	0.0263
C	1	0.4534	0.5007
ACEI_ARB	1	10.7073	0.0011
steroid	1	6.0352	0.0140

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## 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	0.40642	0.20252	4.0273	0.0448	1.501	1.010 2.233
UA_Q4	2	0.52407	0.21016	6.2187	0.0126	1.689	1.119 2.550
UA_Q4	3	0.87149	0.23010	14.3447	0.0002	2.390	1.523 3.753
gender	1	-0.26957	0.14214	3.5967	0.0579	0.764	0.578 1.009
age	1	-0.00682	0.00679	1.0083	0.3153	0.993	0.980 1.007
Proteinuria24h	1	0.00214	0.00959	0.0498	0.8234	1.002	0.983 1.021
GFR_base_CKDEPI	1	-0.00746	0.00286	6.8209	0.0090	0.993	0.987 0.998
MBP	1	-0.00267	0.00518	0.2655	0.6064	0.997	0.987 1.008
M	1	1.31361	0.15261	74.0937	<.0001	3.720	2.758 5.016
E	1	0.47944	0.23133	4.2953	0.0382	1.615	1.026 2.542
S	1	0.35891	0.14181	6.4055	0.0114	1.432	1.084 1.891
T	1	0.32294	0.15354	4.4242	0.0354	1.381	1.022 1.866
T	2	0.73222	0.32158	5.1846	0.0228	2.080	1.107 3.906
C	1	-0.09268	0.13765	0.4534	0.5007	0.911	0.696 1.194
ACEI_ARB	1	-0.46187	0.14115	10.7073	0.0011	0.630	0.478 0.831
steroid	1	-0.32942	0.13409	6.0352	0.0140	0.719	0.553 0.936

## 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	
M	1 M 1
E	1 E 1
S	1 S 1
T	1 T 1
T	2 T 2
C	1 C 1
ACEI_ARB	1 ACEI_ARB 1
steroid	1 steroid 1