

## 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		
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
1482	283	1199	80.90

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## Convergence Status

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

## Model Fit Statistics

Criterion	Without Covariates	With Covariates
-2 LOG L	3674.676	3478.307
AIC	3674.676	3506.307
SBC	3674.676	3557.344

## Testing Global Null Hypothesis: BETA=0

Test	Chi-Square	DF	Pr > ChiSq
Likelihood Ratio	196.3686	14	<.0001
Score	234.1529	14	<.0001
Wald	195.1787	14	<.0001

## Type 3 Tests

Effect	DF	Wald Chi-Square	Pr > ChiSq
UA_Q4	3	12.4306	0.0060
gender	1	3.7879	0.0516
age	1	1.1103	0.2920
Proteinuria24h	1	0.0408	0.8400
GFR_base_CKDEPI	1	6.3641	0.0116
MBP	1	0.8994	0.3429
M	1	84.3965	<.0001
E	1	4.0203	0.0450
S	1	3.9478	0.0469
T	2	8.5389	0.0140
C	1	0.7538	0.3853

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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	0.35897	0.20201	3.1577	0.0756	1.432	0.964 2.127
UA_Q4	2	0.49712	0.20929	5.6419	0.0175	1.644	1.091 2.478
UA_Q4	3	0.79334	0.22693	12.2220	0.0005	2.211	1.417 3.449
gender	1	-0.27486	0.14122	3.7879	0.0516	0.760	0.576 1.002
age	1	-0.00715	0.00679	1.1103	0.2920	0.993	0.980 1.006
Proteinuria24h	1	0.00205	0.01017	0.0408	0.8400	1.002	0.982 1.022
GFR_base_CKDEPI	1	-0.00727	0.00288	6.3641	0.0116	0.993	0.987 0.998
MBP	1	-0.00491	0.00518	0.8994	0.3429	0.995	0.985 1.005
M	1	1.39241	0.15157	84.3965	<.0001	4.025	2.990 5.417
E	1	0.46099	0.22991	4.0203	0.0450	1.586	1.010 2.488
S	1	0.27627	0.13905	3.9478	0.0469	1.318	1.004 1.731
T	1	0.28550	0.15183	3.5360	0.0601	1.330	0.988 1.792
T	2	0.86239	0.31599	7.4484	0.0063	2.369	1.275 4.400
C	1	-0.11673	0.13444	0.7538	0.3853	0.890	0.684 1.158

## 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