The GLMSELECT Procedure

Data Set	DIRS.ALLDATA
Dependent Variable	logtxb
Selection Method	Backward
Select Criterion	Significance Level
Stop Criterion	Significance Level
Stay Significance Level (SLS)	0.05
Effect Hierarchy Enforced	None

Number of Observations Read	1363
Number of Observations Used	1294

Class Level Information						
Class	Levels	Values				
catdose	2	<= 567 mg/wk > 567 mg/wk				
sex	2	Female Male				
smoke	3	Past Mth Past Yr but not past mth Not recently or at all				
lipid	2	Yes No				
diab_comb	2	Yes No				
afibhist	2	Yes No				

Dimensions				
Number of Effects	13			
Number of Parameters	20			

The GLMSELECT Procedure

Backward Selection Summary											
Step	Effect Removed	Number Effects In	Number Parms In	Model R-Square	Adjusted R-Square	AIC	AICC	віс	СР	SBC	PRESS
0		13	14	0.1629	0.1544*	-121.1421*	-120.7665*	-1414.8361*	14.0000*	-1344.8252*	433.3063*

Backward Selection Summary								
Step Removed ASE F Value Pr >								
0		0.3273						

Selection stopped because the next candidate for removal has SLS < 0.05.

Stop Details									
Candidate Compare For Effect Significance Significance									
Removal	afibhist	0.0409	<	0.0500	(SLS)				

The GLMSELECT Procedure **Selected Model**

The selected model is the model at the last step (Step 0).

Effects: Intercept catdose age ratioi egfr logualbumin il6 pselectin sex smoke lipid diab_comb afibhist

Note: The p-values for parameters and effects are not adjusted for the fact that the terms in the model have been selected and so are generally liberal.

Analysis of Variance									
Source DF Sum of Mean Square F Value Pr >									
Model	13	82.42501	6.34039	19.16	<.0001				
Error	1280	423.55832	0.33090						
Corrected Total	1293	505.98333							

0.57524
7.13308
0.1629
0.1544
-121.14212
-120.76653
-1414.83611
14.00000
433.30630
-1344.82521
0.32732

Multiple Regression model predicting LN UTXB Ratio ASA Use = Yes, Adjusting for dose <= 567 vs > 567

The GLMSELECT Procedure Selected Model

Parameter Estimates									
Parameter	DF	Estimate	Standardized Estimate	Standard Error	t Value	Pr > t			
Intercept	1	4.807795	0	0.221479	21.71	<.0001			
catdose <= 567 mg/wk	1	0.226374	0.158261	0.037046	6.11	<.0001			
catdose > 567 mg/wk	0	0	0						
age	1	0.013443	0.179731	0.002282	5.89	<.0001			
ratioi	1	0.000071745	0.068704	0.000028311	2.53	0.0114			
egfr	1	0.008750	0.234936	0.001132	7.73	<.0001			
logualbumin	1	0.056192	0.106786	0.015026	3.74	0.0002			
il6	1	0.014023	0.066116	0.005624	2.49	0.0128			
pselectin	1	0.003081	0.063753	0.001277	2.41	0.0160			
sex Female	1	0.175414	0.139340	0.033437	5.25	<.0001			
sex Male	0	0	0						
smoke Past Mth	1	0.318079	0.125323	0.066739	4.77	<.0001			
smoke Past Yr but not past mth	1	0.050111	0.014682	0.088717	0.56	0.5723			
smoke Not recently or at all	0	0	0						
lipid Yes	1	-0.082829	-0.062817	0.034416	-2.41	0.0162			
lipid No	0	0	0						
diab_comb Yes	1	0.150287	0.097258	0.041444	3.63	0.0003			
diab_comb No	0	0	0						
afibhist Yes	1	0.123742	0.054034	0.060458	2.05	0.0409			
afibhist No	0	0	0						