

JMP[®] Pro 13 Modeling Workflow Enhancements With Predictor Screening, Generalized Regression and the New Formula Depot Karen Copeland – Boulder Statistics

Abstract

- Building models is often an exploratory, iterative process that can result in many saved models. The new Formula Depot in JMP13 Pro provides a workflow to save, compare, and score models without cluttering your data table.
- Prior to building models one may need to screen a large number of factors to identify those with modeling potential. The JMP predictor screening platform is one tool for the job.
- Once a smaller set of factors is identified then the generalized regression personality in the fit model platform is a tool for model building and further factor selection. Generalized regression options include various algorithms and validation options to facilitate factor selection and/or minimize over fitting.
- The generalized regression capabilities have been expanded in JMP Pro 13 to include more types of models, more fitting algorithms and model visualizations, such as ROC curves and profilers.

Predictor Screening

- Predictor screening uses bootstrap forests to identify strong predictors from a pool of predictors.
- Because of the randomness in the procedure, results from re-running the analysis will shift. The larger the set of predictors, the more variability you will see in the results across runs.
- I typically run predictor screening at least 3 times and look at the overall list.

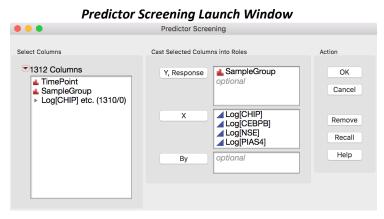
Generalized Regression

- The generalized regression platform is my goto platform for model building.
- Workflow improvements in JMP13 include
 - Re-launch with active effects
 - ROC curves (for categorical responses)

Formula Depot

- The FD is a "container" (a *.jrp file) that holds formulas (remember a model is a formula).
- From many modeling platforms you now have the option to "publish" your model. This saves the prediction formula to the Formula Depot rather than to your data table.
- Once you have a models in the FD you can compare them with the model compare or profiler.
- You can run the model to add it to your data table or to add it to any data table. However, be careful, if your column names or data structures change between tables your model will not run. For example, if you build a model with a gender column coded as M and F but try to run it in a data table with gender as Male and Female the model will not run. Recode is a simple fix.
- For those who need a model outside of JMP you can score the model from the FD. This will translate the JSL code into another coding language (C, Java, Python, SQL, or SAS) for implementation outside of JMP.

Analyze > Screening > Predictor Screening



Y, Repsones = SampleGroup = Gender (10 male, 10 female) X = Predictors = 1310 proteins

> The report window lists the predictors based on their contribution from a bootstrap forest model. I use this as an exploratory tool to help select predictors. I am most interested in the top predictors. The Portion column and corresponding bar chart help to identify how many of the predictors I may wish to explore further.

Predictor Screening Report Window						
Log SL Age Data Example	e - Predictor Screening	of SampleGroup				
🚯 🧞 🖓 🚯	JP / 4	A \ B @				
Window Tools Show Data Table	Graph Tools	Annotation Tools	Column S			
Predictor Screening						
	SampleGroup					
Predictor	Contribution	Portion	Rank			
Log[Myostatin]	0.949746	0.0613	1			
Log[Persephin]	0.562376	0.0363	2			
Log[MOZ]	0.556570	0.0359	3			
Log[LPPL]	0.550751	0.0355	4			
Log[Cytidylate kinase]	0.521042	0.0336	5			
Log[Cystatin M]	0.506290	0.0327	6			
Log[HPV E7 Type 16]	0.482394	0.0311	7			
Log[IL-1 R AcP]	0.435756	0.0281	8			
Log[Troponin T]	0.388162	0.0250	9			
Log[MASP3]	0.382429	0.0247	10			
Log[IL-1 R4]	0.358897	0.0232	11			
Log[PSA]	0.358897	0.0232	12			
Log[TACI]	0.346465	0.0224	13			
Log[Apo A-I]	0.326013	0.0210	14			
Log[GDF-11/8]	0.326013	0.0210	15			
Log[CD22]	0.326013	0.0210	16			
Log[ALK-1]	0.326013	0.0210	17			
Log[OBCAM]	0.326013	0.0210	18			
Log[ANGL4]	0.326013	0.0210	19			
Log[FSH]	0.303164	0.0196	20			
Log[tPA]	0.288412	0.0186	21			
Log[WNT7A]	0.288412	0.0186	22			
	0.277259	0.0179	23			
Log[ASAHL]	0.277259	0.0179	24			
Log[Troponin I, skeletal, fast twitch]	0.277259	0.0179	25			
Log[Luteinizing hormone]	0.273774	0.0177	26			
Log[Thyroxine-Binding Globulin]	0.269921	0.0174	27			
Log[WFKN1]	0.267596	0.0173	28			
Log[TrATPase]	0.267596	0.0173	29			
Log[a1-Antichymotrypsin]	0.257819	0.0166	30			
Log[JAML1]	0.256849	0.0166	31			
Log[3HAO]	0.256849	0.0166	32			
Log[DYRK3]	0.244064	0.0157	33			
	0.000000	0.0154	24			

Analyze > Fit Model : Generalized Regression Personality

	Model Launch Win	dow		
	Fit Model		SL Age Data Example - Gen	eralized R
Model Specification				
elect Columns	Pick Role Variables	Personality: Generalized Regression		Data Tabla
 Ce28 Columns PlatePosition SampleId Barcode TimePoint ExtIdentifier SampleGroup Subject.ID GampleGroupUniqueID CHIP etc. (1310/0) Log[CHIP] etc. (1310/0) 	Y ▲ SampleGroup Weight optional numeric Freq optional numeric Validation optional By optional Construct Model Effects Add Log[App A-1] Log[App A-1] Log[Chr] Log[Chr] Macros + Log[CH1] Attributes ← Log[CH1] Attributes ← No Intercept	Distribution: Binomial Target Level: F Help Run Recall Keep dialog open Remove	Window Tools Show Generalized Regress for SampleGroup = F Model Launch Singularity Details Estimation Method Lasso Adaptive Advanced Control Validation Method AlCc Early Stopping Go 	\$

Tip #1: Predictor Screening report is linked to the data table, highlight column in the report to select in the data table then click Add in the model launch window.

Tip #2: Save the predictor screening results to a data table (right click > Make into Data Table) then copy column names from this table and paste into the model dialog.

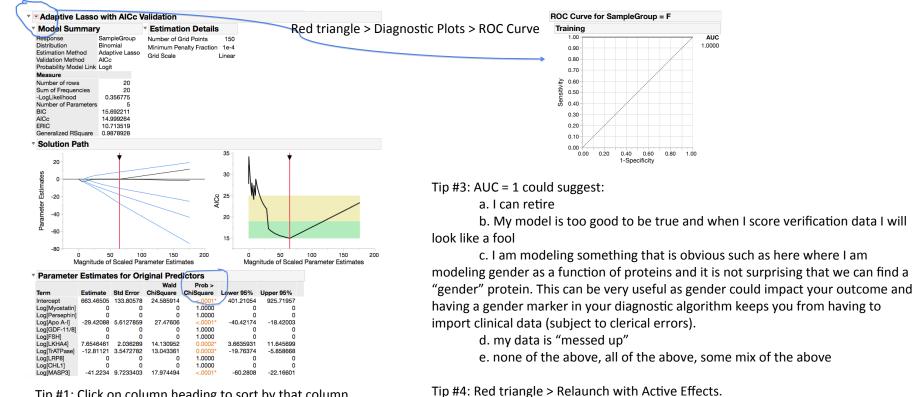
Tip #3: Select columns of interest and then use column reorder to place them together in your data table.

Analyze > Fit Model : Generalized Regression Personality

AUC

1.0000

1.00



Tip #1: Click on column heading to sort by that column. Tip #2: Red triangle > Regression Reports > Active Parameter Estimates.

