

Generalized Linear Models (ST425A)

(Advanced level course, 7.5 hec, Aut. 2019)

Examination (Part 2)

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- **Date and time:** Monday 30 September 2019, 13:00 - 15:00
- **Permitted facilities:** Pocket calculator and attached R- and SAS-codes.
- **Return of exam:** Not yet decided (information will be sent via e-mail or athena).
- **Instructions:**
 - The total amount of points for this part of the exam is 20.
 - The minimum requirement to pass this part of examination is 10 points.
 - Solutions to each question should be detailed enough and well-motivated in order to get full points.

The table below presents the results of a survey on ideal number of children (grouped into 3 increasing categories) among 3663 women cross classified by their education (3 levels) and residence (2 levels).

Education	Residence	Ideal # children		
		≤ 2	3 - 5	> 5
None	Urban	218	200	248
	Rural	731	643	532
Primary	Urban	220	143	121
	Rural	134	44	18
Secondary or above	Urban	250	94	35
	Rural	22	6	4

- a) Fit an appropriate model with *Ideal # children* as response variable and *Education* as explanatory variable.
- b) Repeat (a) with *Education* and *Residence* as explanatory variables.
- c) Use the model-deviances in (a) and (b) to suggest which model is more adequate.
- d) Do the results in (a) and (b) indicate you need to add an interaction term (between *Education* and *Residence*) to the model in (b)?
- e) Interpret your final results on the effect of *Education* and *Residence* on ideal children ever born

Summarize your results in a form of a report that includes choice of a model (with justification), the fitted model, and overall comments on your results (estimates and test statistics). Attach relevant SAS or R codes, tables and figures as appendices.

Generalized Linear Models 2019: R- and SAS Commands

Packages Used
car
ResourceSelection
aod
nnet
MASS
VGAM
vcd
mcprofile
lmtest

Description	Command
Simple/multiple linear regression	lm(formula, data)
One-way / Two-way ANOVA	aov(formula , data)
Type I SS	anova(model)
Type II or III SS	Anova(model , type="II") Anova(model , type="III")
Binary regression	glm(cbind(y,n-y)~ , family=binomial(link="logit"),data) link=c("logit ","probit ","cloglog")
Odds ratio estimates (model with interaction)	predict.glm(model, newdata=data.frame(x1=c(),x2=factor()),se.fit=TRUE)
Multinomial logistic regression	multinom(formula,data,weights)
Proportional odds models	polr(formula,data,weights,Hess=TRUE,method="logistic") method = c("logistic", "probit", "cloglog", "cauchit") vglm(formula=formula, family=cumulative (link="logitlink"),parallel=T, reverse=F), weights,data)
Adjacent-Category model (with proportional odds assumption)	vglm(formula, family=acat (link="loglink", parallel=T, reverse=F), weights,data)

Confidence interval (with assumptions)	<code>vglm(formula, family=ratio(link="loglink", parallel=T, reverse=F), weights,data)</code>	Continuation ratio (with proportional odds assumption)	<code>glm(formula, family=ratio(link="logit", parallel=T, reverse=F))</code>
Rate ratio + CI	<code>exp(confint(mcpfile(model, CM=matrix(C,1))))</code>	Log-linear models	<code>glm(formula, family=Poisson(link="log"), data)</code>
Jack-knife/studentized residuals	<code>rstudent(model)</code>	Standardized residuals	<code>rstANDARD(model)</code>
Cook's distance	<code>cooks.distance(model)</code>	Jack-knife/studentized residuals	<code>rstudent(model)</code>
Leverages	<code>hatvalues(model)</code>	Standarized residuals	<code>rstANDARD(model)</code>
Cook's distance	<code>cooks.distance(model)</code>	Jack-knife/studentized residuals	<code>rstudent(model)</code>
Residuals	<code>residuals(model, type="pearson")</code>	Deviance residuals	<code>residuals(model, type="deviance")</code>
Bayesian information criterion	<code>BIC(model)</code>	Log likelihood	<code>logLik(model)</code>
Akaike's information criterion	<code>AIC(model)</code>	Log likelihood	<code>logLik(model)</code>
Deviance	<code>deviance(model)</code>	Deviance	<code>deviance(model)</code>
Likelihood ratio test	<code>anova(model1, model2, test="Chisq")</code>	Likelihood ratio test	<code>anova(model1, model2, test="Chisq")</code>
Shapiro-Wilk Test	<code>shapiro.test()</code>	Bruegh Pagan Test	<code>bptest(model)</code>
Tests for homogeneity of variance across groups	<code>fligner.test(y~group, data)</code>	Hosmer-Lemeshow's goodness of fit	<code>hoslem.test(y, fitted(model), g=)</code>
Levene's test (y~group, data)	<code>leveneTest(y~group, data)</code>	Hosmer-Lemeshow's goodness of fit	<code>hoslem.test(y, fitted(model), g=)</code>
Wald test	<code>wald.test(b = coef(model), Sigma = vcov(model), Terms =)</code>	Testing for single/joint effects of covariates/factors	<code>linearHypothesis(M1, c("Coef1"=0, "Coef2"=0), test="Chisq")</code>
Linear hypothesis tests (M1, c("Coef1", "Coef2"), test="Chisq")	<code>linearHypothesis(M1, c("Coef1"=0, "Coef2"=0), test="Chisq")</code>	Linear hypothesis tests (M1, c("Coef1", "Coef2"), test="Chisq")	<code>linearHypothesis(M1, c("Coef1"=0, "Coef2"=0), test="Chisq")</code>
Perason chi-Square test for independence	<code>assocstats(H)</code>	Confidence intervals for parameters estimates	<code>confint.default()</code>

Relevel/order factors	relevel(x, ref="...") factor(x, levels=unique(x)) factor(x, ordered=T)
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Description	Command	Simple/multiple linear regression	Binary regression	Multinomial logistic regression	Cumulative logit model	Adjacent-Category model	Continuation ratio (with assumption)	Log-linear models
ANOVA		class A (ref = 'A1'), B (ref = 'B1');						
Descriptive		lsmeans x / adjust = tukey cl;	model formula / link=LOGIT (or PROBIT or CLLOGLOG) aggregate scale = none influence;	proc logistic data = dataset;	output out=tablename pred=logit (or probit or plogit) effectplot fit/obs;	ods graphics on;	ODDSRATIO (model with interaction)	ODDSRATIO name of class var;
Regression		lsmeans x / adjust = tuckey cl;	model formula / link=LOGIT (or PROBIT or CLLOGLOG) aggregate scale = none influence;	proc logistic data = dataset;	output out=tablename pred=logit (or probit or plogit) effectplot fit/obs;	ods graphics off;	freq frequency;	class var (ref="param-ref") var2(ref="param-ref");
Binary regression			model formula / link=LOGIT (or PROBIT or CLLOGLOG) aggregate scale = none influence;	proc logistic data = dataset;	output out=tablename pred=logit (or probit or plogit) effectplot fit/obs;	ods graphics on;	model responses (ref="param-ref") var2(ref="param-ref");	model responses (ref="base-line") var2(link=glogit);
Multinomial logistic regression				ods graphics on;	freq frequency;	run;	oddsratio var;	effectplot interaction (plotby=var1/polybar);
Regression				ods graphics off;	model responses (ref="base-line") var2(link=glogit);	run;	model responses (ref="base-line") var2(link=glogit);	model responses (ref="base-line") var2(link=glogit);
Cumulative logit model					model / LINK = CLOGIT SCALING=none AGGREGATE;	model / LINK = CLOGIT SCALING=none AGGREGATE;	output out=pred predprobs=1 predprobs=c;	output; run;
Adjacent-Category model						continuation ratio (with assumption)	propotional odds (with assumption)	proc genmod data = data;
Continuation ratio (with assumption)						propotional odds (with assumption)	propotional odds (with assumption)	tables var1*var2 / chisq expected nopercent nocol nocum weight n;
Log-linear models								proc freq data = dataset order = data;

			Confidence intervals for Parameter estimates Relative/order factors
			Pearson Chi-Square test for independence Covariates/factors Effects of Testing for single/joint Valid test
			Hosmer-Lemeshow's goodness of fit run: PROC logistic data = data; model formula / link=logit lackfit;
			Tests for homogeneity of variance across groups
			Breusch Pagan Test
			Shapiro-Wilk Test
			Likelihood ratio test
			deviance
			Log likelihood
			Bayesian information criterion
			Akaike's information criterion
			Jackknife/studentized residuals
			Cook's distance
			Leverages
			Pearson residuals
			Deviance residuals
			Standardized residuals
			JACK-knife/studentized residuals
			Rate ratio + CI
			estimate 'RR', var1 1 0 -1 0 var1*var2 1 0...0 /e; xbeta std Reschi Resdev); run;
			ods output obsstats = testl (keep = frequency type site pred model formula / dist = poisson obsstats type3; reference group name"; class var1 (ref = "reference group name") var2 (ref = "

