

# BIST P8110: Applied Regression II

## 19. Intro to PROC GENMOD

Qixuan Chen

Department of Biostatistics  
Columbia University

# PROC GENMOD

- ▶ The GENMOD procedure fits generalized linear models (GLM), such as
  - ▶ linear regression
  - ▶ logistic regression
  - ▶ ordinal logistic regression
  - ▶ Poisson regression
  - ▶ negative binomial regression
  - ▶ zero-inflated Poisson regression
- ▶ The GENMOD procedure can also fit generalized linear models to correlated responses by the GEE method

# SAS Syntax

The following statements are often used in the GENMOD procedure. Items within the < > are optional.

```
PROC GENMOD < options > ;  
  CLASS variables ;  
  MODEL response = < effects > < /options > ;  
  REPEATED SUBJECT = subject-effect < /options > ;  
  RUN ;
```

The REPEATED statement is used in GEE analysis.

# PROC GENMOD Statement

- ▶ The PROC GENMOD statement invokes the procedure.
- ▶ Two most commonly used options are
  - ▶ **DATA**=SAS data-set, to specify the SAS data set containing the data to be analyzed.
  - ▶ **PLOTS**, to specify the plots to be created.

- ▶ Before requesting plots, enable **ODS Graphics**

```
ODS GRAPHICS ON;  
ODS GRAPHICS OFF;
```

- ▶ If to produce all plots available, use options

```
PROC GENMOD PLOTS=all;
```

- ▶ If to request one of the available plots, use options

```
PROC GENMOD PLOTS=PREDICTED;
```

- ▶ The default is to display related multiple plots in a panel. If to display multiple plots individually, use options

```
PROC GENMOD PLOTS (UNPACK) =DFBETA;
```

# CLASS Statement

- ▶ The CLASS statement names the categorical variables to be used in the analysis.
- ▶ The CLASS statement must precede the MODEL statement.
- ▶ The options can be specified for each variable by enclosing them in parentheses after the variable name

```
CLASS treatment (REF='4') age (REF='1') / PARAM = REF ;
```

- ▶ The global options can be also specified for the CLASS statement by placing them after a slash (/).

```
CLASS treatment age / PARAM = REF REF =first;
```

# MODEL Statement

- ▶ The MODEL statement specifies the response and the explanatory variables.

```
MODEL response = <effects> </options> ;
```

- ▶ The **response** can be specified in the form of a single variable or in the form of a ratio of two variables denoted **events/trials**.
  - ▶ The first form is applicable to all responses. The second form is applicable only to grouped binomial response data.
- ▶ The **options** often use
  - ▶ **COVB**, **DIST**, **LINK**, **OFFSET**, **SCALE**, and **TYPE3**.

## MODEL Statement Options

- ▶ COVB – requests that the parameter estimate covariance matrix be displayed.
- ▶ TYPE3 – requests that statistics for Type 3 contrasts be computed for each effect specified in the MODEL statement.
- ▶ DIST – specifies the built-in probability distribution to use in the model.

DIST=	Distribution
NOR	normal
BIN	binomial
MULT	multinomial
POI	Poisson
NB	negative binomial
ZIP	zero-inflated Poisson
ZINB	zero-inflated negative binomial

# MODEL Statement Options

- ▶ LINK – specifies the link function to be used in the model.

LINK=	Link Function
ID	identity
LOG	log
LOGIT	logit

- ▶ OFFSET – specifies a variable in the input data set to be used as an offset variable.
- ▶ SCALE – specifies an overdispersed model.
  - ▶ EX: In Poisson distribution, we specify  $\text{var}(Y) = \phi\lambda$ . The dispersion parameter  $\phi$  can be estimated using Pearson's chi-square statistic or Deviance divided by the degrees of freedom

```
MODEL y = x / SCALE=P;
```

```
MODEL y = x / SCALE=D;
```