

STAT 705 Chapter 18: ANOVA diagnostics and remedial measures

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Stat 705: Data Analysis II

18.1 Residuals

Raw residual is

$$e_{ij} = Y_{ij} - \bar{Y}_{i\bullet} = Y_{ij} - \hat{Y}_{ij}.$$

Studentized residual is

$$r_{ij} = \frac{e_{ij}}{se\{e_{ij}\}} = \frac{e_{ij}}{\sqrt{MSE(1 - h_{ij})}} = \frac{e_{ij}}{\sqrt{MSE(n_i - 1)/n_i}}.$$

Studentized deleted residual is

$$t_{ij} = \frac{e_{ij}}{\sqrt{MSE_{(ij)}(1 - h_{ij})}}.$$

Recall if model is correct then

$$t_{ij} \sim t(n_T - r - 1).$$

Question: Is the model reasonable?

Use residuals to check (p.778):

- 1 Constant error variance.
- 2 Outliers.
- 3 Normality.

To check these,

- 1 SAS default graphics plots e_{ij} vs. \hat{Y}_{ij} . Should show roughly constant spread.
- 2 SAS plots t_{ij} vs. \hat{Y}_{ij} . Can formally determine if ij th observation is outlier by checking

$$|t_{ij}| > t\left(1 - \frac{\alpha}{2n_T}; n_T - r - 1\right).$$

- 3 Normal probability plot of $\{e_{ij}\}$ should be reasonably straight.

Also, if data are collected over time, can plot e_{ij} vs time observation was recorded to check for serial correlation.

```
data kenton;
input sales design @@;
datalines;
  11 1 17 1 16 1 14 1 15 1 12 2 10 2 15 2 19 2 11 2
  23 3 20 3 18 3 17 3 27 4 33 4 22 4 26 4 28 4
;

proc glm data=kenton plots=all; class design;
  model sales=design;
run;
```

18.2 Tests for constant variance

Expanded model is

$$Y_{ij} \stackrel{ind.}{\sim} N(\mu_i, \sigma_i^2),$$

want to test $H_0 : \sigma_1 = \dots = \sigma_r$.

In `proc glm` use `means factorname / hovtest`; “hov” stands for homogeneity of variance. The options are `hovtest=Bartlett` (not robust to non-normality), `hovtest=Levene` (Default), `hovtest=Obrien`, and `hovtest=BF` (Brown-Forsythe, pp. 784–785).

Levene's test is simple oneway ANOVA F-test on “dispersion variables” $z_{ij}^2 = (Y_{ij} - \bar{Y}_{i\bullet})^2$ (default) or $z_{ij} = |Y_{ij} - \bar{Y}_{i\bullet}|$ (`type=abs`). O'Brien and Brown-Forsythe simply use different z_{ij} – Brown-Forsythe uses $z_{ij} = |Y_{ij} - \tilde{Y}_{i\bullet}|$, also called the *modified Levene test*.

Five different types of flux, response is amount of force (lbs) required to break soldered joint.

```
data abt;
input force flux @@;
datalines;
  14.87 1 16.81 1 15.83 1 15.47 1 13.60 1 14.76 1 17.40 1 14.62 1
  18.43 2 18.76 2 20.12 2 19.11 2 19.81 2 18.43 2 17.16 2 16.40 2
  16.95 3 12.28 3 12.00 3 13.18 3 14.99 3 15.76 3 19.35 3 15.52 3
   8.59 4 10.90 4  8.60 4 10.13 4 10.28 4  9.98 4  9.41 4 10.04 4
  11.55 5 13.36 5 13.64 5 12.16 5 11.62 5 12.39 5 12.05 5 11.95 5
;

proc glm data=abt plots=all;
class flux;
model force=flux;
means flux / hovtest=bf;
run;
```

18.3 Remedial measures

- Weighted least squares for non-constant variance, but otherwise normal data (Section 18.4). Alternatively, add `welch` in a `means` statement in `proc glm` to perform an ANOVA generalization of Satterthwaite's two-sample approach.
- Transformation to normality, e.g. Box-Cox for non-normal data with non-constant variance (Section 18.5). Carried out the same way as in STAT 704.
- A nonparametric test (Section 18.7), e.g. the Kruskal-Wallis test. Available in the `npar1way` procedure in SAS. Doesn't care about normality or constant variance. Essentially replaces the Y_{ij} with ranks obtained from ranking all observations without regard to group and computes regular ANOVA F-test.

Servo-Data, Inc., operates mainframes at three locations, all identical make/model. Recorded are lengths of time (hours) between failures at the three locations.

```
data servo;
input time loc @@;
datalines;
    4.41 1 100.65 1 14.45 1 47.13 1 85.21 1
    8.24 2 81.16 2 7.35 2 12.29 2 1.61 2
    106.19 3 33.83 3 78.88 3 342.81 3 44.33 3
;

* look at diagnostic plots;
proc glm data=servo plots=all; class loc;
model time=loc;
means loc;
run;

* Box-Cox transformation;
proc transreg data=servo;
model boxcox(time) = class(loc);
run;

* Kruskal-Wallis test;
proc npar1way data=servo;
class loc; var time;
run;
```

18.4 Weighted least squares

Special case of Section 11.1 (pp. 421–427). Define weights $w_{ij} = 1/s_i^2$ where s_i is sample standard deviation from i th factor level.

Tim prefers to fit the model $Y_{ij} = \mu_i + \epsilon_{ij}$, $\epsilon_{ij} \stackrel{ind.}{\sim} N(0, \sigma_i^2)$ directly in `proc mixed`. This approach uses maximum likelihood and normal approximations for testing. However, WLS also uses an approximation. `proc mixed` will at least take variability of estimating σ_i by s_i into account!

Carried out in `proc mixed` by adding

```
repeated / group=factorname;
```

Carried out in `proc glimmix` by adding

```
random _residual_ / group=factorname;
```

Fitting

$$Y_{ij} = \mu + \alpha_i + \epsilon_{ij}, \quad \epsilon_{ij} \stackrel{ind.}{\sim} N(0, \sigma_i^2).$$

```
data abt;
input force flux @@;
datalines;
  14.87 1 16.81 1 15.83 1 15.47 1 13.60 1 14.76 1 17.40 1 14.62 1
  18.43 2 18.76 2 20.12 2 19.11 2 19.81 2 18.43 2 17.16 2 16.40 2
  16.95 3 12.28 3 12.00 3 13.18 3 14.99 3 15.76 3 19.35 3 15.52 3
   8.59 4 10.90 4  8.60 4 10.13 4 10.28 4  9.98 4  9.41 4 10.04 4
  11.55 5 13.36 5 13.64 5 12.16 5 11.62 5 12.39 5 12.05 5 11.95 5
;

proc sort data=abt; by flux; run;

proc mixed data=abt;
class flux;
model force=flux / solution;
repeated / group=flux; * different variance for each flux;
run;

* another way using Welch-Satterthwaite approximation;
proc glm data=abt; class flux;
model force=flux;
means flux / welch;
run;
```

The “Null Model Likelihood Ratio Test” tests

$H_0 : \sigma_1 = \sigma_2 = \sigma_3 = \sigma_4 = \sigma_5$ by comparing two models, the model with different factor level variances, and the model with one overall variance.

The Type III test looks at $H_0 : \alpha_1 = \alpha_2 = \alpha_3 = \alpha_4 = \alpha_5 = 0$, whether there are significant differences in group means. This is of course the same as testing $H_0 : \mu_1 = \mu_2 = \mu_3 = \mu_4 = \mu_5$ in the cell-means model.

What happens if you run the usual one-way ANOVA?

According to a Minitab White Paper by Rob Kelly:

- **Normality:** Simulations show accuracy with non-normal data with reasonably large sample sizes. If r is 2–9 levels, need $n_i \geq 15$. If r is 10–12 levels, need $n_i \geq 20$.
- **Non-constant variance.** The Type I error is best for equal group sizes: true α ranges from 0.02 – 0.08 when $\alpha = 0.05$. Otherwise α goes as high as 0.22 in some cases.

If you have *many* comparisons to make, check out `proc multtest`. Among other techniques, `multtest` can implement the false discovery rate control approach in Benjamini and Hochberg (1995).

What about multiple comparisons?

The Games-Howell (Games and Howell, 1996) method has been shown to work very well for normal data with non-constant variance and different groups sizes as long as $n_i \geq 5$. Simulation can also work well.

```
proc mixed data=abt; class flux;
  model force=flux / ddfm=satterth; * denom. degrees freedom;
  repeated / group=flux;          * Games-Howell adjust;
  lsmeans flux / adjust=smm adjdfe=row;
run;
```

```
proc glimmix data=abt; class flux;
  model force=flux;
  random _residual_ / group=flux;
  covtest homogeneity;
  lsestimate flux "1 vs 2" -1 1 0 0 0,
                 "3 vs 4" 0 0 -1 1 0 / adjust=simulate;
run;
```

Can also follow up Kruskal-Wallis with two-sample Mann-Whitney-Wilcoxin tests, using Bonferroni to bound the FER. Need to have moderate to small r to achieve any power. A better approach is to use the `dscf` option in `proc npar1way` to compute the Dwass, Steel, Critchlow-Fligner multiple comparison analysis, based on pairwise two-sample MWW comparisons.