

# Analysis of Variance Illustration – Stata version 14

*Dear BIOSTATS 640 Spring 2023  
Please be aware that this illustration may be out of date.  
Thank you - cb*

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## 1. Preliminaries

Preliminary – Download the stata data set **hers\_640anova.dta**.

**Note – This data set is accessible through the internet. Alternatively, you can download it from the course website.**

(a) In Stata, input directly from the internet using the command [use](#)  
use “[http://people.umass.edu/biep640w/datasets/hers\\_640anova.dta](http://people.umass.edu/biep640w/datasets/hers_640anova.dta)”, clear

(b) From the course website, right click to download. Afterwards, in Stata, use **FILE > OPEN**  
See, <http://people.umass.edu/biep640w/webpages/demonstrations.html>

Preliminary – Download the stata command **anovaplot**

. \* Download "add-on" anova command anovaplot if you don't already have it  
. findit anovaplot

Preliminary – Download the stata command **anovacontrast**

. \* Download "add-on" anova command anovacontrast if you don't already have it  
. findit anovacontrast

## 2. Illustration: Trial of Estrogen and Progesterone for Prevention of Heart Disease

### Source

Hulley et al (1998) Randomized trial of estrogen plus progestin for secondary prevention of heart disease in postmenopausal women. The Heart and Estrogen/progestin Replacement Study. *Journal of the American Medical Association*, **280**(7), 605-613

### Source Data

The actual data set contains information on  $n=2,763$ . This was a randomized controlled trial investigation of hormone therapy for the prevention of heart attack and death.

### The Data for this Illustration is a subset

- (1) This dataset is a subset of size  $n=612$  from a larger data set of Hulley et al (1998) that is described below. I have taken a subsample of size  $n=612$  so that students with “small stata” will be able to reproduce this analysis.
- (2) Specifically, I took a random sample of 300 whites, all 218 African-Americans, plus all 94 women of other race ethnicity.

### Analysis Question

In this hypothetical data set, does systolic blood pressure (**sbp**) vary by race-ethnicity (**raceth**)?

### 3. ONE WAY Analysis of Variance

#### One Way Analysis of Variance Question

In this hypothetical data set, does  $Y$  = systolic blood pressure (**sbp**) vary over groups defined by  $X$ =race-ethnicity (**raceth**)?

#### 3a. Descriptives – Graphs

```
. ** Set scheme for graphs - user choice
. set scheme s1color

. * get min and max of y=sbp for y-axis tick marks
. tabstat sbp, stat(min max)
```

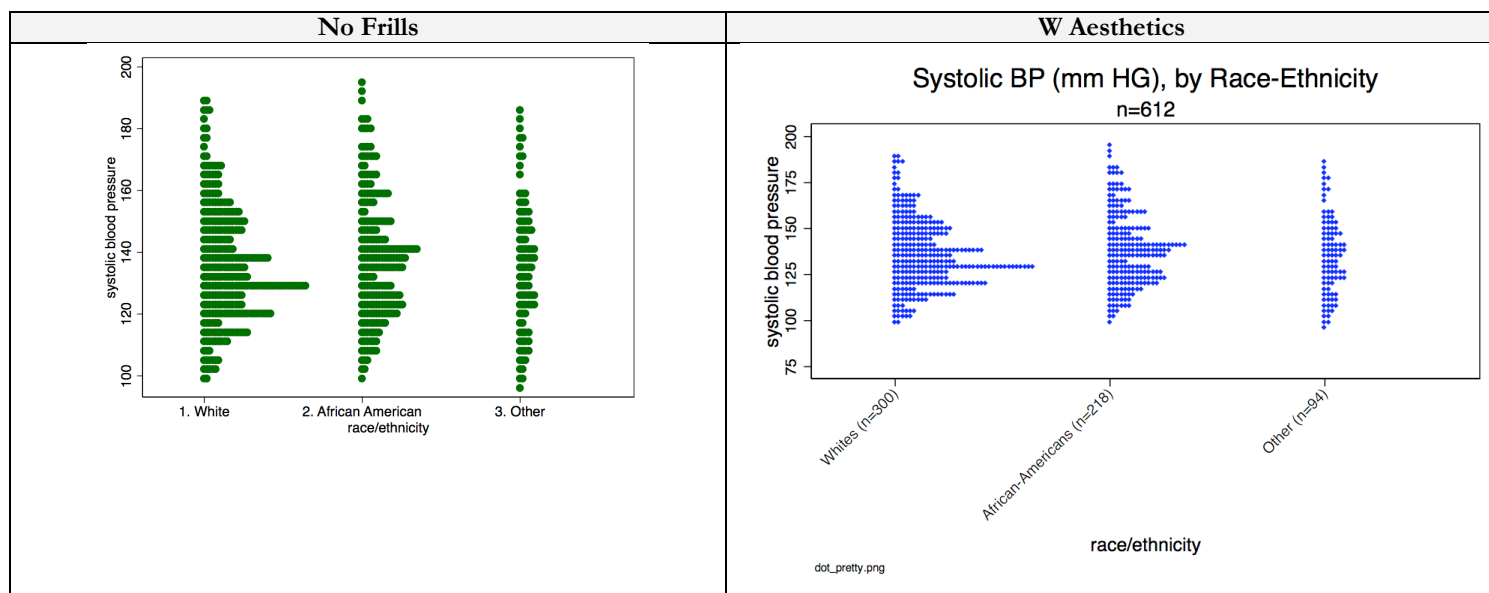
variable	min	max
-----+-----		
sbp	95	194
-----+-----		

```
. * retrieve correspondence between codes and code labels and get sample sizes
. numlabel, add
. tabulate raceth
```

race/ethnicity	Freq.	Percent	Cum.
-----+-----			
1. White	300	49.02	49.02
2. African American	218	35.62	84.64
3. Other	94	15.36	100.00
-----+-----			
Total	612	100.00	

```
. ** Side-by-side dot plot - NO FRILLS
. sort raceth
. dotplot sbp, over(raceth)
```

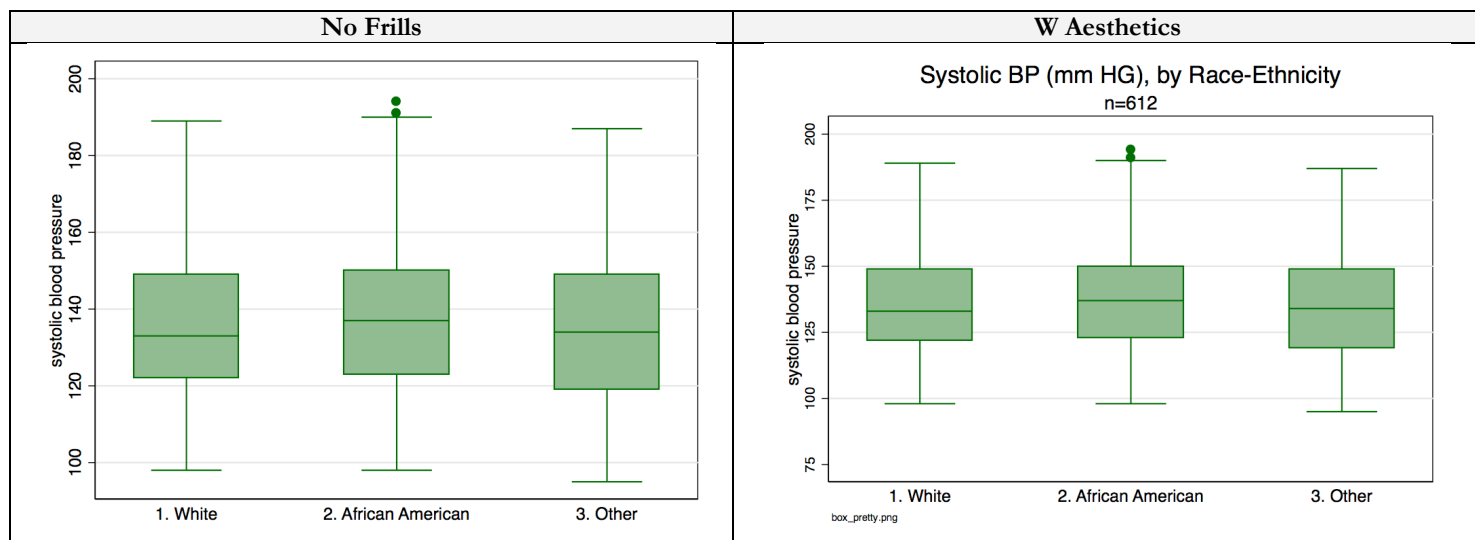
```
. ** Side-by-side dot plot - w AESTHETICS
. dotplot sbp, over(raceth) msymbol(d) msize(vsmall) mcolor(blue) ylabel(75(25)200, labsize(small)) xlabel(1 "Whites (n=300)" 2 "African-Americans (n=218)" 3 "Other (n=94)", labsize(small) angle(45)) title("Systolic BP (mm HG), by Race-Ethnicity")
subtitle("n=612") caption("dot_pretty.png", size(vsmall))
```



*sbp distributions are similar in the 3 groups. It's unlikely that anova will be significant.*

```
. ** Side-by-side box plot - NO FRILLS
. graph box sbp, over(raceth)

. ** Side-by-side box plot - w AESTHETICS
. graph box sbp, over(raceth) ylabel(75(25)200, labsize(small)) title("Systolic BP (mm HG), by Race-Ethnicity") subtitle("n=612")
caption("box_pretty.png", size(vsmall))
```



### 3b. Descriptives – Numerical

Note - There is more than one way to do these!

```
. * Descriptives - Numerical Descriptives of Raw Data
. tabstat sbp, by(raceth) stat(n mean sd sem min q max)
```

Summary for variables: sbp  
by categories of: raceth (race/ethnicity)

raceth	N	mean	sd	se(mean)	min	p25	p50	p75	max
1. White	300	136.0133	18.55138	1.071064	98	122	133	149	189
2. African Ameri	218	138.2339	19.99252	1.354064	98	123	137	150	194
3. Other	94	135.1809	21.25977	2.192778	95	119	134	149	187
Total	612	136.6765	19.50878	.7885956	95	122	135	149	194

```
. tabulate raceth, summarize(sbp)
```

race/ethnicity	Mean	Std. Dev.	Freq.
White	136.01333	18.551379	300
African A	138.23394	19.992518	218
Other	135.18085	21.259767	94
Total	136.67647	19.508777	612

Numerical descriptives confirm what we saw in the graphs. The means are similar. We see that they range 135.2 mm Hg to 138.2 mm Hg. The standard deviations appear to be similar, too. They range 18.55 mm Hg to 21.26 mm Hg. Test of equality of variances is likely to be non-significant.

```
. * Descriptives - Obtain group specific means and associated 95% confidence intervals
. sort raceth
. by raceth: ci means sbp
```

```
-> raceth = White
```

Variable	Obs	Mean	Std. Err.	[95% Conf. Interval]
sbp	300	136.0133	1.071064	133.9056 138.1211

```
-> raceth = African American
```

Variable	Obs	Mean	Std. Err.	[95% Conf. Interval]
sbp	218	138.2339	1.354064	135.5651 140.9027

```
-> raceth = Other
```

Variable	Obs	Mean	Std. Err.	[95% Conf. Interval]
sbp	94	135.1809	2.192778	130.8264 139.5353

No surprise. Lots of overlap of the confidence intervals.

### 3c. Analysis of Variance Model Estimation

*Stata offers at least 2 commands for a one way anova: oneway or anova.*

#### KEY:

If you use **oneway**, then the predictor variable is allowed to be **string** or **numeric**

If you use **anova**, then the predictor variable **must be numeric**.

**How to convert a string variable into a new, numeric variable.**

Use the command **encode** with the option **generate** as follows:

**encode *stringvar*, generate(*numericvar*)**

Tip – STATA will automatically create variable value labels for your new numericvar.

**Now you have two choices for performing your one way anova fit:**

(1) **oneway *yvariable factor***

(2) **anova *yvariable numericfactor***

```
.* Illustration of command ANOVA
.* anova YVARIABLE NUMERICFACTOR
. anova sbp raceth
```

```
Number of obs = 612      R-squared      = 0.0037
Root MSE      = 19.5042  Adj R-squared = 0.0005
```

Source	Partial SS	df	MS	F	Prob > F
Model	871.000171	2	435.500085	1.14	0.3190
raceth	871.000171	2	435.500085	1.14	0.3190
Residual	231670.941	609	380.412054		
Total	232541.941	611	380.592375		

*The null hypothesis assumption of equal means does not lead to an unlikely outcome ( $p=.32$ ); the null hypothesis is NOT rejected.*

```

.* TIP - ONEWAY provides Bartlett test of equal variances.
.* oneway YVARIABLE FACTOR
. oneway sbp raceth

```

Source	Analysis of Variance				
	SS	df	MS	F	Prob > F
Between groups	871.000171	2	435.500085	1.14	0.3190
Within groups	231670.941	609	380.412054		
Total	232541.941	611	380.592375		

Bartlett's test for equal variances:  $\chi^2(2) = 3.1766$  Prob> $\chi^2 = 0.204$

```

.* To obtain summary statistics, use command ONEWAY with option TABULATE
. oneway sbp raceth, tabulate

```

race/ethnicity	Summary of systolic blood pressure		
	Mean	Std. Dev.	Freq.
1. White	136.01333	18.551379	300
2. Africa	138.23394	19.992518	218
3. Other	135.18085	21.259767	94
Total	136.67647	19.508777	612

Source	Analysis of Variance				
	SS	df	MS	F	Prob > F
Between groups	871.000171	2	435.500085	1.14	0.3190
Within groups	231670.941	609	380.412054		
Total	232541.941	611	380.592375		

Bartlett's test for equal variances:  $\chi^2(2) = 3.1766$  Prob> $\chi^2 = 0.204$

```

.* To obtain reference cell coding regression solution, execute 2 commands
.* First - anova
.* Second - regress

```

```

.* Step 1 - Command is anova
. anova sbp raceth

```

Number of obs = 612  
Root MSE = 19.5042

R-squared = 0.0037  
Adj R-squared = 0.0005

Source	Partial SS	df	MS	F	Prob > F
Model	871.000171	2	435.500085	1.14	0.3190
raceth	871.000171	2	435.500085	1.14	0.3190
Residual	231670.941	609	380.412054		
Total	232541.941	611	380.592375		



```
.* Step 2 - Command is regress
. regress
```

Source	SS	df	MS	Number of obs	=	612
Model	871.000171	2	435.500085	F(2, 609)	=	1.14
Residual	231670.941	609	380.412054	Prob > F	=	0.3190
				R-squared	=	0.0037
				Adj R-squared	=	0.0005
Total	232541.941	611	380.592375	Root MSE	=	19.504

	sbp	Coef.	Std. Err.	t	P> t	[95% Conf. Interval]
	raceth					
African American		2.220612	1.735814	1.28	0.201	-1.188296 5.629519
Other		-.8324823	2.305423	-0.36	0.718	-5.360027 3.695063
	_cons	136.0133	1.126073	120.79	0.000	133.8019 138.2248

Checking with means on page 8 – they match!

$$\hat{Y}_{\text{WHITE}} = \hat{Y}_{\text{ref}} = [\hat{b}_0] = 136.01$$

$$\hat{Y}_{\text{AFRICAN-AMERICAN}} = \hat{Y}_2 = [\hat{b}_0 + \hat{b}_2] = [136.01 + 2.22] = 138.23$$

$$\hat{Y}_{\text{OTHER}} = \hat{Y}_3 = [\hat{b}_0 + \hat{b}_3] = [136.01 - 0.83] = 135.18$$

### 3d. Post-hoc Pairwise Comparison of Groups

*Pairwise comparisons of groups is done using the command `pwcompare`.*

*Note – You must have fit the model first using `anova`*

```
.* PRELIMINARY - Must first fit model using anova
. anova sbp raceth
```

Number of obs = 612					
R-squared = 0.0037					
Root MSE = 19.5042					
Adj R-squared = 0.0005					
Source	Partial SS	df	MS	F	Prob > F
Model	871.000171	2	435.500085	1.14	0.3190
raceth	871.000171	2	435.500085	1.14	0.3190
Residual	231670.941	609	380.412054		
Total	232541.941	611	380.592375		

```
.* Compare pairwise means with NO adjustment for multiple comparisons using PWCOMPARE
.* pwcompare FACTOR
. pwcompare raceth
```

Pairwise comparisons of marginal linear predictions

Margins : asbalanced

	Contrast	Std. Err.	Unadjusted [95% Conf. Interval]	
raceth				
2 vs 1	.9654558	1.034007	-1.065196	2.996108
3 vs 1	2.228419	1.373318	-.4685942	4.925432
3 vs 2	1.262963	1.433615	-1.552466	4.078392

*For all pairwise comparisons of groups, the 95% confidence interval includes the null hypothesis value of zero.*

```
. * Pairwise comparison of means using Bonferroni adjustment (NOT RECOMMENDED)
. pwcompare raceth, mcompare(bonferroni) sort effects
```

Pairwise comparisons of marginal linear predictions

Margins : asbalanced

	Number of Comparisons
raceth	3

	Contrast	Std. Err.	Bonferroni t	P> t	Bonferroni [95% Conf. Interval]
raceth					
3 vs 2	-3.053094	2.406646	-1.27	0.615	-8.830518 2.724331
3 vs 1	-.8324823	2.305423	-0.36	1.000	-6.36691 4.701945
2 vs 1	2.220612	1.735814	1.28	0.604	-1.946404 6.387628

*Even with the stringent Bonferroni adjustment, for all pairwise comparisons of groups, no statistically significant differences are found. Not surprising, given what we've already seen.*

```
. * Pairwise comparison of means using Tukey adjustment for Multiple Comparisons
. * NOTE - This requires equal sample sizes in all groups
. * So, technically, I should not have done this.
. pwcompare raceth, mcompare(tukey)
```

Pairwise comparisons of marginal linear predictions

Margins : asbalanced

	Number of Comparisons
raceth	3

	Contrast	Std. Err.	Tukey [95% Conf. Interval]
raceth			
2 vs 1	2.220612	1.735814	-1.857604 6.298827
3 vs 1	-.8324823	2.305423	-6.24897 4.584005
3 vs 2	-3.053094	2.406646	-8.7074 2.601212

Note: The tukey method requires balanced data for proper level coverage. A factor was found to be unbalanced.

*Yes, yes, I know. I should not have done the Tukey procedure*

3e. Post-Hoc Graphs

PRELIMINARY – You must have fit the model first using anova

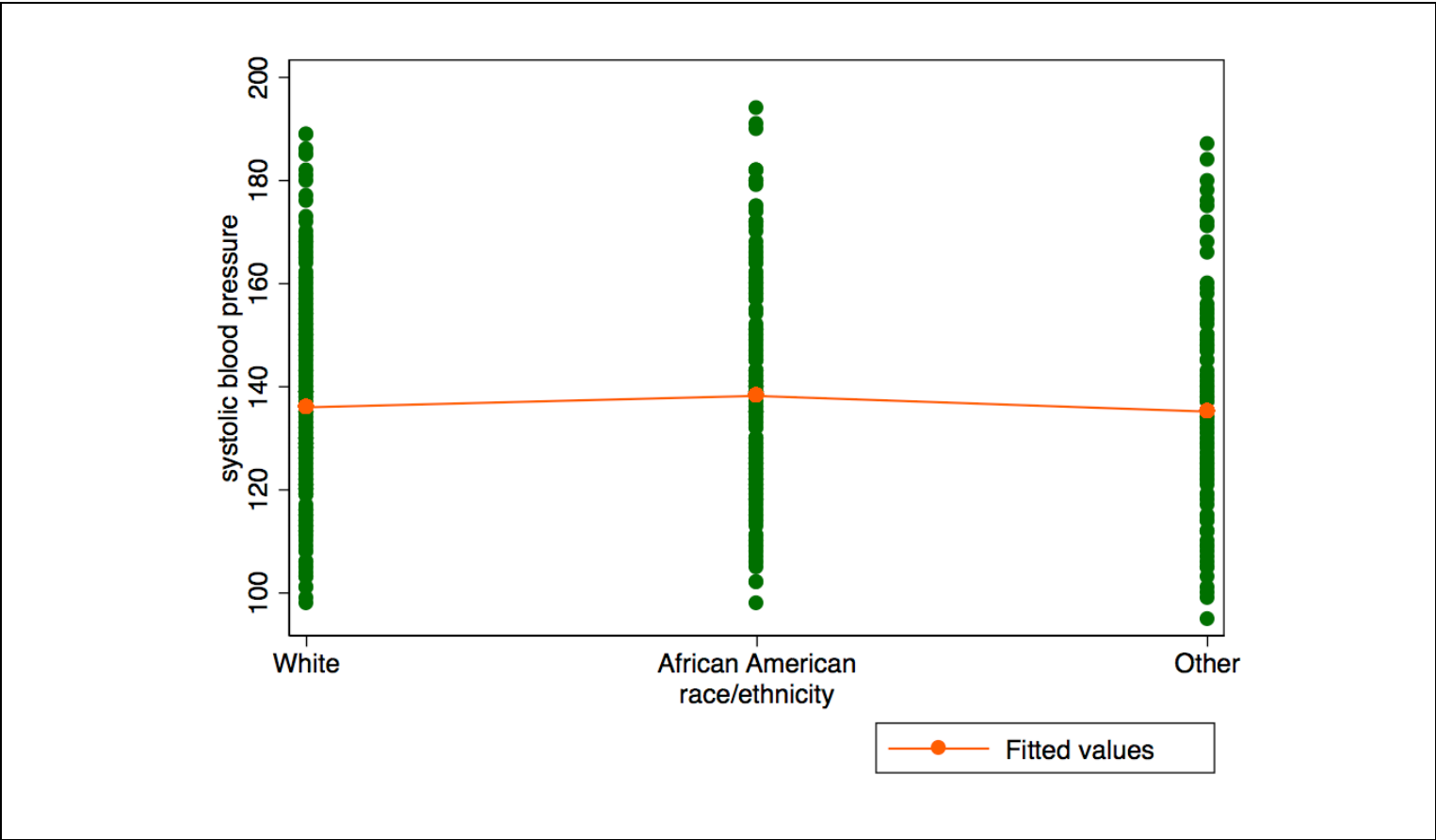
```
. anova sbp raceth
```

Number of obs = 612  
Root MSE = 19.5042

R-squared = 0.0037  
Adj R-squared = 0.0005

Source	Partial SS	df	MS	F	Prob > F
Model	871.000171	2	435.500085	1.14	0.3190
raceth	871.000171	2	435.500085	1.14	0.3190
Residual	231670.941	609	380.412054		
Total	232541.941	611	380.592375		

```
. * anova plot - NO FRILLS  
. anovaplot
```



## 4. ONE WAY Analysis of Variance Diagnostics

### 4a. Homogeneity of Variances

. \* BARTLETT's Test is provided in output from command oneway  
 . \* Caution: This test is sensitive to the assumption of normality  
 . oneway sbp raceth

Source	Analysis of Variance			F	Prob > F
	SS	df	MS		
Between groups	871.000171	2	435.500085	1.14	0.3190
Within groups	231670.941	609	380.412054		
Total	232541.941	611	380.592375		

Bartlett's test for equal variances:  $\chi^2(2) = 3.1766$  Prob> $\chi^2 = 0.204$

*The null hypothesis of equal variances is not rejected (Bartlett test p-value=.20)*

. \* LEVENE and BROWN-FORSYTHE tests are obtained using the command robvar  
 . \* These are good choices when assumption of normality is in question.  
 . \* W\_0 = Levene test  
 . \* W\_50 = Forsythe-Browne modification of Levene test (mean is replaced by median)  
 . \* W\_10 = Forsythe-Browne modification of Levene test (mean is replaced by 10% trim)  
 . \* robvar(YVAR), by(FACTOR)  
 . robvar sbp, by(raceth)

race/ethnicity	Summary of systolic blood pressure		
	Mean	Std. Dev.	Freq.
White	136.01333	18.551379	300
African A	138.23394	19.992518	218
Other	135.18085	21.259767	94
Total	136.67647	19.508777	612

W0 = 1.4143305 df(2, 609) Pr > F = 0.24388559 ← Levene  
 W50 = 1.4701779 df(2, 609) Pr > F = 0.23069929 ← Brown-Forsythe with median  
 W10 = 1.4741613 df(2, 609) Pr > F = 0.22978655 ← Brown-Forsythe with 10% trimmed mean

*The null hypothesis of equal variances is not rejected by Levene's test either (p-value=.24)*

## 4b. Normality of Residuals

*NOTE – The following requires that you fit your ONE WAY anova model using the command `anova`. The following will not work if you fit your model using the command `oneway`.*

```
. anova sbp raceth
```

Number of obs = 612					
R-squared = 0.0037					
Root MSE = 19.5042					
Adj R-squared = 0.0005					
Source	Partial SS	df	MS	F	Prob>F
Model	871.00017	2	435.50009	1.14	0.3190
raceth	871.00017	2	435.50009	1.14	0.3190
Residual	231670.94	609	380.41205		
Total	232541.94	611	380.59238		

```
. predict ehat, residuals
```

```
. * Shapiro-Wilk Test (NULL: Distribution is normal)
```

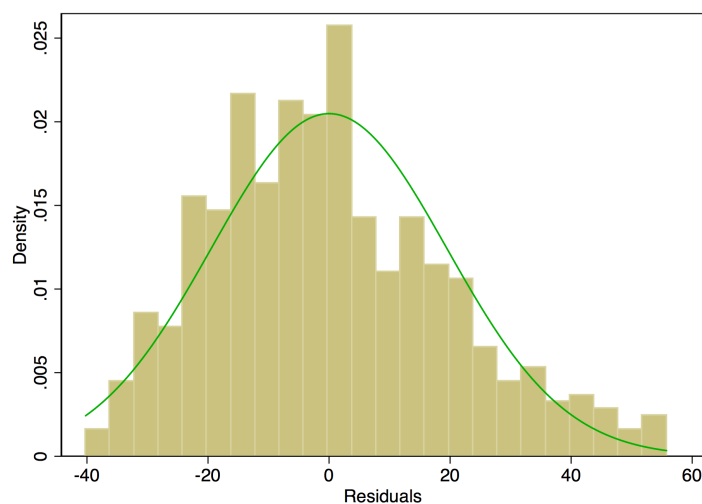
```
. swilk ehat
```

Shapiro-Wilk W test for normal data

Variable	Obs	W	V	z	Prob>z
ehat	612	0.98066	7.810	4.986	0.00000

*Boo hoo. There is statistically significant evidence of departure from normality. Null is rejected ( $p < .00001$ ). Let's do a graphical look to see if we're really in trouble.*

```
. histogram ehat, normal
```



*This departure from normality is not so severe as to warrant transforming the data. For purposes of this illustration, we'll proceed under the assumption of normality.*

## 5. TWO WAY Analysis of Variance

### Two Way Analysis of Variance Question

In this hypothetical data set, does  $Y$  = systolic blood pressure (**sbp**) vary over groups defined by  $X_1$ =race-ethnicity (**raceth**) and  $X_2$ =physical activity (**physact**)?

#### 5a. Descriptives – Graphs

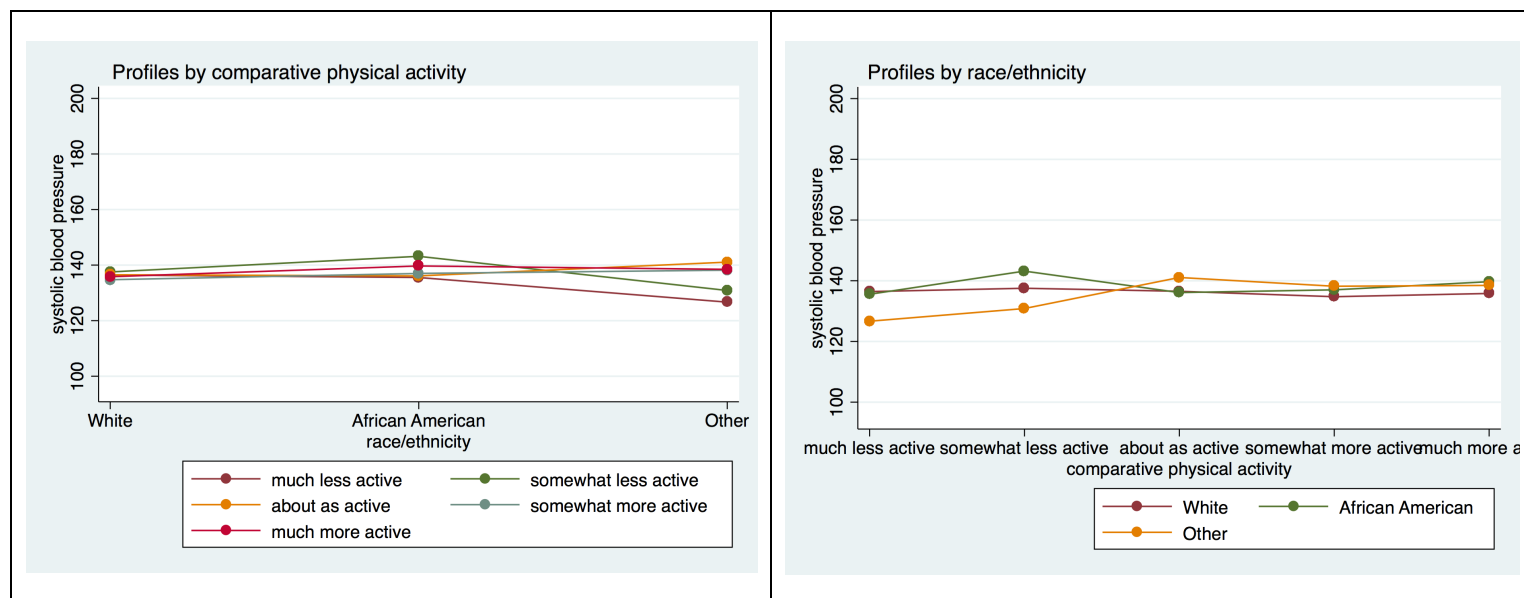
We will use the command **anovaplot** here.

Our goal at this point is just a visualization of the means to see if there is interaction present. However, the stata command **anovaplot** requires that we have first executed the command **anova**. So we will do this “quietly,” which means that the output will not be shown.

```
. * Obtain plot of mean Y=sbp over groups in 2 ways to assess interaction
. quietly: anova sbp raceth physact raceth#physact

.* Plot of means of Y=sbp with raceth, SEPARATELY for groups defined by physact (left panel)
. anovaplot raceth physact, scatter(msym(none))

.* Plot of means of Y=sbp with physact, SEPARATELY for groups defined by raceth (right panel)
. anovaplot physact raceth, scatter(msym(none))
```



*Okay, maybe this was not the best choice of example. These mean profiles look pretty similar.*

## 5b. Descriptives – Numeric

```
. table physact raceth, contents(n sbp mean sbp sd sbp)
```

comparative physical activity	race/ethnicity		
	White	African American	Other
much less active	14	33	18
	136.4286	135.5455	126.6667
	19.63793	18.88467	22.68454
somewhat less active	55	51	21
	137.5273	143.1765	130.8571
	18.71012	21.08621	19.1371
about as active	99	67	26
	136.5253	136.1045	141.0769
	17.62208	20.27921	21.07875
somewhat more active	105	44	16
	134.7333	137	138.1875
	18.94337	19.55196	19.73565
much more active	27	23	13
	135.8148	139.6956	138.4615
	20.47895	18.5948	22.59283

*Pretty tidy summary! A quick scan, however, suggests that the means are not likely to be assessed as statistically significantly different. Not surprising given the graphs we obtained earlier.*

## 5c. Analysis of Variance Model Estimation

```
. anova sbp raceth physact raceth#physact
```

Number of obs =		612	R-squared =	0.0258	
Root MSE =		19.4804	Adj R-squared =	0.0029	
Source	Partial SS	df	MS	F	Prob>F
Model	5988.6916	14	427.76368	1.13	0.3300
raceth	762.56725	2	381.28362	1.00	0.3668
physact	1149.1915	4	287.29788	0.76	0.5535
raceth#physact	3563.4604	8	445.43255	1.17	0.3125
Residual	226553.25	597	379.48618		
Total	232541.94	611	380.59238		

*Oh dear. How boring. Nothing significant here.*





```
. * Post-hoc Tests of Simple Main Effects
. * PRELIMINARY - Must have fit model w anova command first
. anova sbp raceth physact raceth#physact
```

```
Number of obs =      612    R-squared      = 0.0258
Root MSE      =    19.4804    Adj R-squared = 0.0029
```

Source	Partial SS	df	MS	F	Prob>F
Model	5988.6916	14	427.76368	1.13	0.3300
raceth	762.56725	2	381.28362	1.00	0.3668
physact	1149.1915	4	287.29788	0.76	0.5535
raceth#physact	3563.4604	8	445.43255	1.17	0.3125
Residual	226553.25	597	379.48618		
Total	232541.94	611	380.59238		

```
. * Test of simple main effect of predictor = physact, separately at each raceth
. sme physact raceth
```

```
Test of physact at raceth(1): F(4/597) = .21576196
Test of physact at raceth(2): F(4/597) = 1.2545496
Test of physact at raceth(3): F(4/597) = 1.9011494
```

Critical value of F for alpha = .05 using ...

```
-----
Dunn's procedure          = 2.9791402
Marascuilo & Levin       = 3.2193716
per family error rate    = 3.0491547
simultaneous test procedure = 4.8816539
```

*At each level of raceth, the F-statistic for testing equality of sbp over groups defined by physact is NOT in the critical region. Thus, we have no statistically significant evidence of variations in sbp by physact.*

```
. * Test of simple main effect of predictor = raceth, separately at each physact
. sme raceth physact
```

```
Test of raceth at physact(1): F(2/597) = 1.4432739
Test of raceth at physact(2): F(2/597) = 3.1481138
Test of raceth at physact(3): F(2/597) = .66958053
Test of raceth at physact(4): F(2/597) = .35753139
Test of raceth at physact(5): F(2/597) = .25655016
```

Critical value of F for alpha = .05 using ...

```
-----
Dunn's procedure          = 4.003167
Marascuilo & Levin       = 4.414349
per family error rate    = 4.6408772
simultaneous test procedure = 9.7633078
```

*At each level of physact, the F-statistic for testing equality of sbp over groups defined by raceth is NOT in the critical region. Thus, we have no statistically significant evidence of variations in sbp by raceeth*

```
. * Pairwise comparison of groups using command anovacontrast (requires installation)
. * CAUTION!!!! If interaction is significant, this analysis may not be appropriate.
```

```
. * Preliminary: Use command fre to see again the codings for raceth
. fre raceth
```

```
raceth -- race/ethnicity
```

		Freq.	Percent	Valid	Cum.
Valid	1 White	300	49.02	49.02	49.02
	2 African American	218	35.62	35.62	84.64
	3 Other	94	15.36	15.36	100.00
	Total	612	100.00	100.00	

```
. * Pairwise comparison of 2 groups: (1=White) v (2=African American)
. * KEY: Entries in option values( ) must total 0 and must make sense
. * (1=White) → 1, (2=African American) → -1 and (3=Other) → 0
. quietly: anova sbp raceth physact raceth#physact
. anovacontrast raceth, values(1 -1 0)
```

Contrast variable: raceth (1 -1 0) Dep Var: sbp

source	SS	df	MS	Contrast =	
				N of obs =	612
contrast	397.018529	1	397.0185	F =	1.05
error	226553.25	597	379.4862	Prob > F =	0.3068
				t =	1.02

Overall, there is no statistically significant difference in sbp, Whites v African-Americans ( $p=.31$ )

```
. * Another example
. fre physact
```

```
physact -- comparative physical activity
```

		Freq.	Percent	Valid	Cum.
Valid	1 much less active	65	10.62	10.62	10.62
	2 somewhat less active	127	20.75	20.75	31.37
	3 about as active	192	31.37	31.37	62.75
	4 somewhat more active	165	26.96	26.96	89.71
	5 much more active	63	10.29	10.29	100.00
	Total	612	100.00	100.00	

```
. * Compare all "less" active versus all "more" active
. quietly: anova sbp raceth physact raceth#physact
```

```
. anovacontrast physact, values(-1 -1 0 1 1)
```

Contrast variable: physact (-1 -1 0 1 1) Dep Var: sbp

source	SS	df	MS	Contrast =	
				N of obs =	612
contrast	378.775904	1	378.7759	F =	1.00
error	226553.25	597	379.4862	Prob > F =	0.3182
				t =	1.00

Overall, there is no statistically significant difference in sbp, Less active v More active ( $p=.32$ )

## 6. TWO WAY Analysis of Variance Diagnostics

### 6a. Homogeneity of Variances

Spoiler – we need to trick Stata.

We will use the command **robvar** to obtain our test of equality of variance. However, STATA will not accommodate groups defined by 2 factors (in our case *raceth* and *physact*). So we will use the command **egen** with the option **group** to create a new variable that is just one collection of groups. In this case, I've named my new variable *allgroups*.

```
. egen allgroups=group(raceth physact)
. robvar sbp, by(allgroups)
```

group(raceth physact)	Summary of systolic blood pressure		
	Mean	Std. Dev.	Freq.
1	136.42857	19.637931	14
2	137.52727	18.710123	55
3	136.52525	17.622078	99
4	134.73333	18.94337	105
5	135.81481	20.478952	27
6	135.54545	18.884668	33
7	143.17647	21.08621	51
8	136.10448	20.279213	67
9	137	19.551958	44
10	139.69565	18.594806	23
11	126.66667	22.684537	18
12	130.85714	19.137099	21
13	141.07692	21.078753	26
14	138.1875	19.735649	16
15	138.46154	22.592828	13
Total	136.67647	19.508777	612

W0 = 0.27440108 df(14, 597) Pr > F = 0.99621084

W50 = 0.24684554 df(14, 597) Pr > F = 0.99786588

W10 = 0.27125653 df(14, 597) Pr > F = 0.99643754

*NOT Significant. We have no statistically significant evidence of a departure from equality of variances*

## 6b. Normality of Residuals

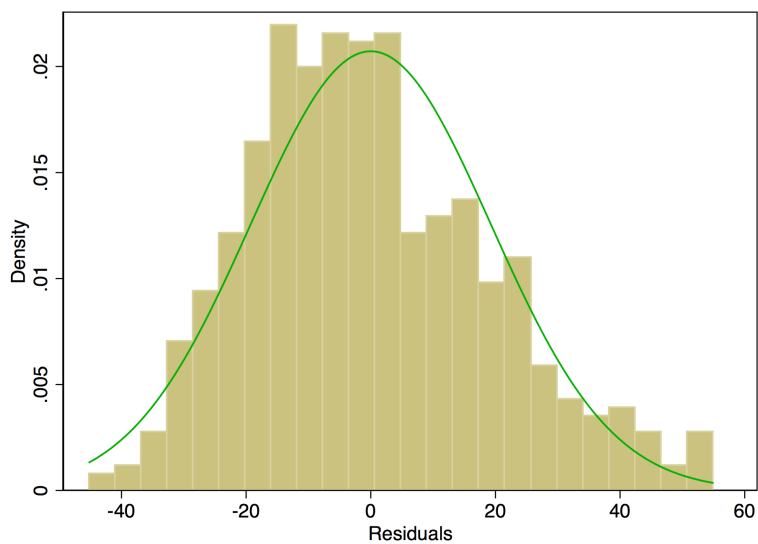
*Same as before(see again Section 4.b)*

```
. quietly: anova sbp raceth physact raceth#physact
. predict ehat, residuals
. * Shapiro-Wilk Test (NULL: Distribution is normal)
. swilk ehat
```

Shapiro-Wilk W test for normal data

Variable	Obs	W	V	z	Prob>z
-----+-----					
ehat	612	0.98058	7.840	4.995	0.00000

```
. histogram ehat, normal
```



*This departure from normality is not so horrible.*