

STAT 401A - Statistical Methods for Research Workers

Two-way ANOVA

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This is also referred to as a **full factorial** or **fully crossed** design.

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For all of these questions, we want to know

- is there any effect and
- if yes, what is the nature of the effect.

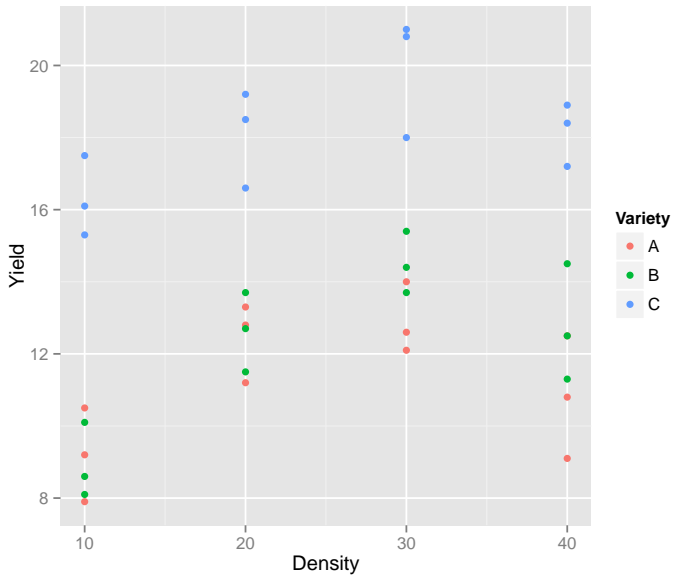
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- is there any effect and
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Confidence intervals can answer these questions.



Summary statistics

Number of replicates

	Variety	10	20	30	40
1	A	3	3	3	3
2	B	3	3	3	3
3	C	3	3	3	3

Mean Yield

	Variety	10	20	30	40
1	A	9.200000	12.43333	12.90000	10.80000
2	B	8.933333	12.63333	14.50000	12.76667
3	C	16.300000	18.10000	19.93333	18.16667

Standard deviation of yield

	Variety	10	20	30	40
1	A	1.300000	1.096966	0.9848858	1.7000000
2	B	1.040833	1.101514	0.8544004	1.6165808
3	C	1.113553	1.345362	1.6772994	0.8736895

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	10	20	30	40
A	μ_{11}	μ_{12}	μ_{13}	μ_{14}
B	μ_{21}	μ_{22}	μ_{23}	μ_{24}
C	μ_{31}	μ_{32}	μ_{33}	μ_{34}

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- 4 The additive model:

$$\begin{aligned}\mu_i = & \beta_0 + \beta_1 I(V_i = A) + \beta_2 I(V_i = B) \\ & + \beta_3 I(D_i = 10) + \beta_4 I(D_i = 20) + \beta_5 I(D_i = 30).\end{aligned}$$

β_1 is the expected difference in yield between varieties A and C at any fixed density

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$$\mu_i = \beta_0 + \beta_1 I(V_i = A) + \beta_2 I(V_i = B) \\ + \beta_3 I(D_i = 10) + \beta_4 I(D_i = 20) + \beta_5 I(D_i = 30).$$

β_1 is the expected difference in yield between varieties A and C at any fixed density

- ⑤ The cell-means model:

$$\mu_i = \beta_0 + \beta_1 I(V_i = A) + \beta_2 I(V_i = B) \\ + \beta_3 I(D_i = 10) + \beta_4 I(D_i = 20) + \beta_5 I(D_i = 30) \\ + \beta_6 I(V_i = A)I(D_i = 10) + \beta_7 I(V_i = A)I(D_i = 20) + \beta_8 I(V_i = A)I(D_i = 30) \\ + \beta_9 I(V_i = B)I(D_i = 10) + \beta_{10} I(V_i = B)I(D_i = 20) + \beta_{11} I(V_i = B)I(D_i = 30)$$

β_1 is the expected difference in yield between varieties A and C at a density of 40

ANOVA Table

ANOVA Table - Additive model

Source	SS	df	MS	F
Factor A	SSA	(I-1)	SSA/(I-1)	MSA/MSE
Factor B	SSB	(J-1)	SSB/(J-1)	MSB/MSE
Error	SSE	n-I-J+1	SSE/(n-I-J+1)	
Total	SST	n-1		

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Factor B	SSB	(J-1)	$SSB/(J-1)$	MSB/MSE
Error	SSE	$n-I-J+1$	$SSE/(n-I-J+1)$	
Total	SST	$n-1$		

ANOVA Table - Cell-means model

Source	SS	df	MS	
Factor A	SSA	I-1	$SSA/(I-1)$	MSA/MSE
Factor B	SSB	J-1	$SSB/(J-1)$	MSB/MSE
Interaction AB	SSAB	$(I-1)(J-1)$	$SSAB / (I-1)(J-1)$	$MSAB/MSE$
Error	SSE	$n-IJ$	$SSE/(n-IJ)$	
Total	SST	$n-1$		

Additive vs cell-means

Opinions differ on whether to use an additive vs a cell-means model when the interaction is not significant.

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Interpretation	Direct	Complicated
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We will continue using the cell-means model to answer the scientific questions of interest.

Two-way ANOVA using PROC GLM

```
DATA tomato;  
  INFILE 'Ch13-tomato.csv' DSD FIRSTOBS=2;  
  INPUT variety $ density yield;  
  
PROC GLM DATA=tomato PLOTS=all;  
  CLASS variety density;  
  MODEL yield = variety|density / SOLUTION;  
  LSMEANS variety / cl adjust=tukey;  
  LSMEANS density / cl adjust=tukey;  
  LSMEANS variety*density / cl adjust=tukey;  
RUN;
```

Two-way ANOVA using PROC GLM

The GLM Procedure

Dependent Variable: yield

Source	DF	Sum of Squares	Mean Square	F Value	Pr > F
Model	11	422.3155556	38.3923232	24.22	<.0001
Error	24	38.0400000	1.5850000		
Corrected Total	35	460.3555556			

R-Square	Coeff Var	Root MSE	yield Mean
0.917368	9.064568	1.258968	13.88889

Source	DF	Type I SS	Mean Square	F Value	Pr > F
variety	2	327.5972222	163.7986111	103.34	<.0001
density	3	86.6866667	28.8955556	18.23	<.0001
variety*density	6	8.0316667	1.3386111	0.84	0.5484

Source	DF	Type III SS	Mean Square	F Value	Pr > F
variety	2	327.5972222	163.7986111	103.34	<.0001
density	3	86.6866667	28.8955556	18.23	<.0001
variety*density	6	8.0316667	1.3386111	0.84	0.5484

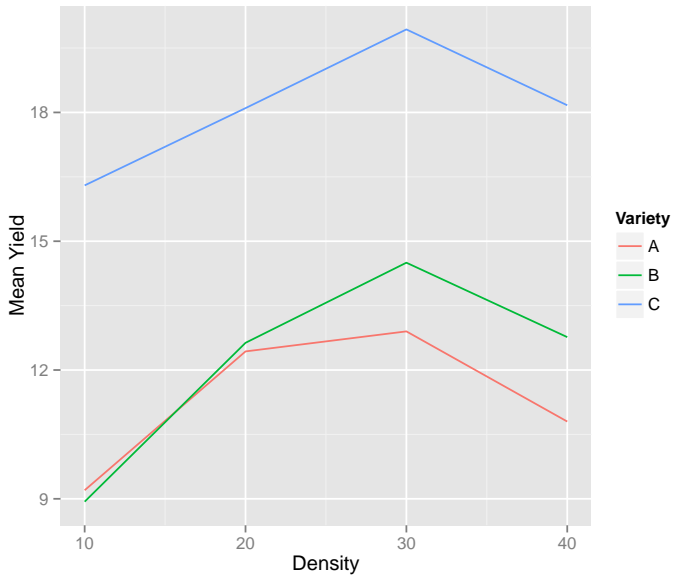
The Type I and Type III SS are equal because the design is balanced.

Two-way ANOVA using PROC GLM

```
MODEL yield = variety|density / SOLUTION;
```

The GLM Procedure

Parameter		Estimate	Standard Error	t Value	Pr > t
Intercept		18.16666667 B	0.72686542	24.99	<.0001
variety	A	-7.36666667 B	1.02794293	-7.17	<.0001
variety	B	-5.40000000 B	1.02794293	-5.25	<.0001
variety	C	0.00000000 B	.	.	.
density	10	-1.86666667 B	1.02794293	-1.82	0.0819
density	20	-0.06666667 B	1.02794293	-0.06	0.9488
density	30	1.76666667 B	1.02794293	1.72	0.0986
density	40	0.00000000 B	.	.	.
variety*density	A 10	0.26666667 B	1.45373083	0.18	0.8560
variety*density	A 20	1.70000000 B	1.45373083	1.17	0.2537
variety*density	A 30	0.33333333 B	1.45373083	0.23	0.8206
variety*density	A 40	0.00000000 B	.	.	.
variety*density	B 10	-1.96666667 B	1.45373083	-1.35	0.1887
variety*density	B 20	-0.06666667 B	1.45373083	-0.05	0.9638
variety*density	B 30	-0.03333333 B	1.45373083	-0.02	0.9819
variety*density	B 40	0.00000000 B	.	.	.
variety*density	C 10	0.00000000 B	.	.	.



Is the mean yield for variety A different from B on average?

```
LSMEANS variety / cl adjust=tukey;
```

Least Squares Means
Adjustment for Multiple Comparisons: Tukey

...

Least Squares Means for effect variety
Pr > |t| for H0: LSMean(i)=LSMean(j)

Dependent Variable: yield

i/j	1	2	3
1		0.2249	<.0001
2	0.2249		<.0001
3	<.0001	<.0001	

variety	yield LSMEAN	95% Confidence Limits	
A	11.333333	10.583245	12.083422
B	12.208333	11.458245	12.958422
C	18.125000	17.374912	18.875088

Least Squares Means for Effect variety

		Difference	Simultaneous 95%	
		Between	Confidence Limits for	
i	j	Means	LSMean(i)-LSMean(j)	
1	2	-0.875000	-2.158534	0.408534
1	3	-6.791667	-8.075201	-5.508132
2	3	-5.916667	-7.200201	-4.633132

Is the mean yield at density 10 different from density 20 on average?

```
LSMEANS density / cl adjust=tukey;
```

Least Squares Means
Adjustment for Multiple Comparisons: Tukey
...

density	yield LSMEAN	95% Confidence Limits	
10	11.477778	10.611650	12.343905
20	14.388889	13.522762	15.255016
30	15.777778	14.911650	16.643905
40	13.911111	13.044984	14.777238

Least Squares Means for Effect density				
		Difference	Simultaneous 95%	
		Between	Confidence Limits for	
i	j	Means	LSMean(i)-LSMean(j)	
1	2	-2.911111	-4.548299	-1.273923
1	3	-4.300000	-5.937188	-2.662812
1	4	-2.433333	-4.070521	-0.796145
2	3	-1.388889	-3.026077	0.248299
2	4	0.477778	-1.159410	2.114966
3	4	1.866667	0.229479	3.503855

Is mean yield different for particular combinations?

```
LSMEANS variety*density / cl adjust=tukey;
```

variety	density	yield LSMEAN	95% Confidence Limits	
A	10	9.200000	7.699824	10.700176
A	20	12.433333	10.933157	13.933510
A	30	12.900000	11.399824	14.400176
A	40	10.800000	9.299824	12.300176
B	10	8.933333	7.433157	10.433510
B	20	12.633333	11.133157	14.133510
B	30	14.500000	12.999824	16.000176
B	40	12.766667	11.266490	14.266843
C	10	16.300000	14.799824	17.800176
C	20	18.100000	16.599824	19.600176
C	30	19.933333	18.433157	21.433510
C	40	18.166667	16.666490	19.666843

Is mean yield different for particular combinations?

```
LSMEANS variety*density / cl adjust=tukey;
```

Least Squares Means for Effect variety*density

		Difference	Simultaneous 95%	
		Between	Confidence Limits for	
i	j	Means	LSMean(i)-LSMean(j)	
1	2	-3.233333	-6.939704	0.473037
1	3	-3.700000	-7.406371	0.006371
1	4	-1.600000	-5.306371	2.106371
1	5	0.266667	-3.439704	3.973037
1	6	-3.433333	-7.139704	0.273037
1	7	-5.300000	-9.006371	-1.593629
1	8	-3.566667	-7.273037	0.139704
1	9	-7.100000	-10.806371	-3.393629
1	10	-8.900000	-12.606371	-5.193629
1	11	-10.733333	-14.439704	-7.026963
1	12	-8.966667	-12.673037	-5.260296
2	3	-0.466667	-4.173037	3.239704
2	4	1.633333	-2.073037	5.339704
2	5	3.500000	-0.206371	7.206371
2	6	-0.200000	-3.906371	3.506371
2	7	-2.066667	-5.773037	1.639704
2	8	-0.333333	-4.039704	3.373037
2	9	-3.866667	-7.573037	-0.160296
2	10	-5.666667	-9.373037	-1.960296
2	11	-7.500000	-11.206371	-3.793629
2	12	-5.733333	-9.439704	-2.026963
3	4	2.100000	-1.606371	5.806371
3	5	3.966667	0.260296	7.673037
3	6	0.266667	-3.439704	3.973037

```
tomato = read.csv("Ch13-tomato.csv")
tomato$Density = factor(tomato$Density)
m = lm(Yield~Variety*Density, tomato)
anova(m)
```

Analysis of Variance Table

Response: Yield

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
Variety	2	327.60	163.799	103.3430	1.608e-12 ***
Density	3	86.69	28.896	18.2306	2.212e-06 ***
Variety:Density	6	8.03	1.339	0.8445	0.5484
Residuals	24	38.04	1.585		

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```
library(lsmmeans)
lsmeans(m, pairwise~Variety)
```

```
$lsmeans
  Variety    lsmean      SE df lower.CL upper.CL
A          11.33333 0.3634327 24 10.58325 12.08342
B          12.20833 0.3634327 24 11.45825 12.95842
C          18.12500 0.3634327 24 17.37491 18.87509
```

Results are averaged over the levels of: Density
Confidence level used: 0.95

```
$contrasts
  contrast estimate      SE df t.ratio p.value
A - B      -0.875000 0.5139715 24  -1.702  0.2249
A - C      -6.791667 0.5139715 24 -13.214 <.0001
B - C      -5.916667 0.5139715 24 -11.512 <.0001
```

Results are averaged over the levels of: Density
P value adjustment: tukey method for a family of 3 means

```
lsmeans(m, pairwise~Density)
```

```
$lsmeans
```

Density	lsmean	SE	df	lower.CL	upper.CL
10	11.47778	0.4196559	24	10.61165	12.34391
20	14.38889	0.4196559	24	13.52276	15.25502
30	15.77778	0.4196559	24	14.91165	16.64391
40	13.91111	0.4196559	24	13.04498	14.77724

Results are averaged over the levels of: Variety
Confidence level used: 0.95

```
$contrasts
```

contrast	estimate	SE	df	t.ratio	p.value
10 - 20	-2.9111111	0.5934831	24	-4.905	0.0003
10 - 30	-4.3000000	0.5934831	24	-7.245	<.0001
10 - 40	-2.4333333	0.5934831	24	-4.100	0.0022
20 - 30	-1.3888889	0.5934831	24	-2.340	0.1169
20 - 40	0.4777778	0.5934831	24	0.805	0.8514
30 - 40	1.8666667	0.5934831	24	3.145	0.0213

Results are averaged over the levels of: Variety
P value adjustment: tukey method for a family of 4 means

```
lsmeans(m, pairwise~Variety*Density)
```

```
$lsmeans
```

Variety	Density	lsmean	SE	df	lower.CL	upper.CL
A	10	9.200000	0.7268654	24	7.699824	10.70018
B	10	8.933333	0.7268654	24	7.433157	10.43351
C	10	16.300000	0.7268654	24	14.799824	17.80018
A	20	12.433333	0.7268654	24	10.933157	13.93351
B	20	12.633333	0.7268654	24	11.133157	14.13351
C	20	18.100000	0.7268654	24	16.599824	19.60018
A	30	12.900000	0.7268654	24	11.399824	14.40018
B	30	14.500000	0.7268654	24	12.999824	16.00018
C	30	19.933333	0.7268654	24	18.433157	21.43351
A	40	10.800000	0.7268654	24	9.299824	12.30018
B	40	12.766667	0.7268654	24	11.266490	14.26684
C	40	18.166667	0.7268654	24	16.666490	19.66684

Confidence level used: 0.95

```
$contrasts
```

contrast	estimate	SE	df	t.ratio	p.value
A,10 - B,10	0.2666667	1.027943	24	0.259	1.0000
A,10 - C,10	-7.1000000	1.027943	24	-6.907	<.0001
A,10 - A,20	-3.2333333	1.027943	24	-3.145	0.1284
A,10 - B,20	-3.4333333	1.027943	24	-3.340	0.0873
A,10 - C,20	-8.9000000	1.027943	24	-8.658	<.0001
A,10 - A,30	-3.7000000	1.027943	24	-3.599	0.0507
A,10 - B,30	-5.3000000	1.027943	24	-5.156	0.0013
A,10 - C,30	-10.7333333	1.027943	24	-10.442	<.0001
A,10 - A,40	-1.6000000	1.027943	24	-1.557	0.9085
A,10 - B,40	-3.5666667	1.027943	24	-3.470	0.0668
A,10 - C,40	-8.9666667	1.027943	24	-8.723	<.0001
B,10 - C,10	-7.3666667	1.027943	24	-7.166	<.0001
B,10 - A,20	-3.5000000	1.027943	24	-3.405	0.0764
B,10 - B,20	-2.7000000	1.027943	24	-2.599	0.0507
B,10 - C,20	-6.1666667	1.027943	24	-5.985	<.0001
B,10 - A,30	-2.5000000	1.027943	24	-2.433	0.0229
B,10 - B,30	-1.8000000	1.027943	24	-1.746	0.0913
B,10 - C,30	-5.2666667	1.027943	24	-5.128	<.0001
B,10 - A,40	-0.9000000	1.027943	24	-0.877	0.3874
B,10 - B,40	0.0000000	1.027943	24	0.000	1.0000
B,10 - C,40	-4.4000000	1.027943	24	-4.281	<.0001
C,10 - C,20	2.0000000	1.027943	24	1.943	0.0642
C,10 - C,30	2.6000000	1.027943	24	2.528	0.0181
C,10 - C,40	1.8000000	1.027943	24	1.746	0.0913
C,20 - C,30	1.6000000	1.027943	24	1.557	0.9085
C,20 - C,40	0.0000000	1.027943	24	0.000	1.0000
C,30 - C,40	1.0000000	1.027943	24	0.973	0.3367

Summary

- Use LSMEANS to answer questions of scientific interest.

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- Check model assumptions
- Consider alternative models, e.g. treating density as continuous