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```
clear;
```

9.1-2

```
control = [17.9 12.2 14.9 13.8 26.1 15.4 20.3 16.9 20.8 14.8];
deficient = [ 7.0 6.9 13.3 11.1 11.0 16.5 12.7 12.4 17.1 9.0];
slow = [19.8 20.3 16.1 17.9 12.4 12.5 17.4 19.9 27.3 14.4];

data = [ control deficient slow ];
groups(1:10) = {'c'};
groups(11:20) = {'d'};
groups(21:30) = {'s'};

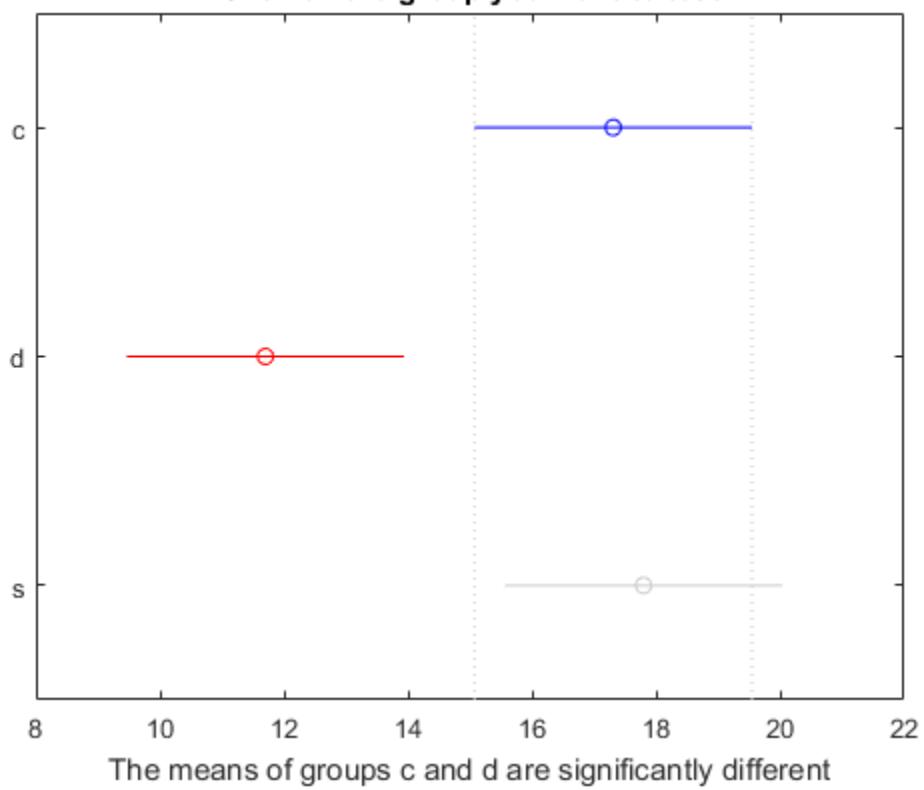
[p,table,stats] = anoval(data,groups)
multcompare(stats);
% yes, heights differ

p =
    0.0034
table =
    'Source'      'SS'          'df'        'MS'        'F'
    'Prob>F'
    'Groups'      [229.7407]    [ 2]      [114.8703]    [7.0587]
[0.0034]
    'Error'       [439.3890]    [27]      [ 16.2737]    []
    []
    'Total'       [669.1297]    [29]      []          []
    []
stats =
    gnames: {3x1 cell}
    n: [10 10 10]
    source: 'anolval'
    means: [17.3100 11.7000 17.8000]
    df: 27
    s: 4.0341
```

ANOVA Table

| Source | SS | df | MS | F | Prob>F |
|--------|---------|----|--------|------|--------|
| <hr/> | | | | | |
| Groups | 229.741 | 2 | 114.87 | 7.06 | 0.0034 |
| Error | 439.389 | 27 | 16.274 | | |
| Total | 669.13 | 29 | | | |

Click on the group you want to test



9.1-4

```
clear;
g0 = [12.7 14.1 13.2];
g05 = [13.5 14.5 14.6];
g1 = [12.7 13.4 13.2];
g2 =[12.7 13.6 14.1];
g3 = [13.4 13.5 14.3];
g4 = [14.5 13.5 14.9];

data = [g0,g05,g1,g2,g3,g4];
groups(1:3) = {'0 g'};
groups(4:6) = {'0.5 g'};
groups(7:9) = {'1 g'};
groups(10:12) = {'2 g'};
groups(13:15) = {'3 g'};
groups(14:18) = {'4 g'};

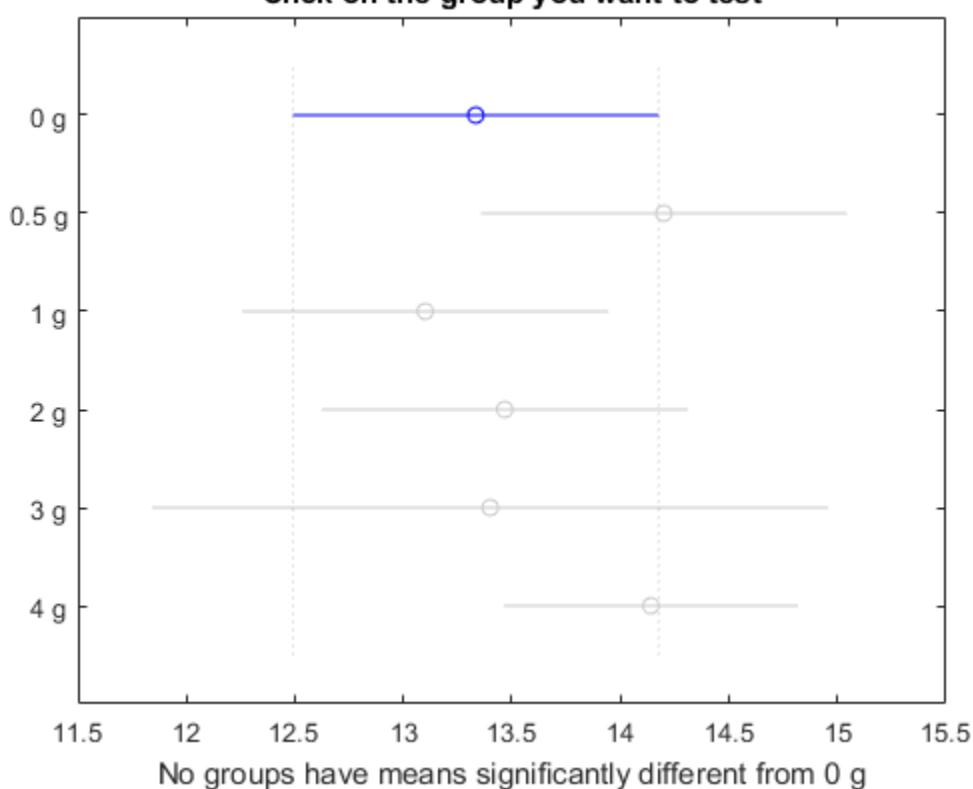
[p,table,stats] = anoval(data,groups)
multcompare(stats);
% no differences

p =
    0.1843
table =
    'Source'      'SS'        'df'        'MS'        'F'        'Prob>F'
    'Groups'      [ 3.4524]   [ 5]       [ 0.6905]   [ 1.8150]   [ 0.1843]
    'Error'       [ 4.5653]   [ 12]      [ 0.3804]   [ ]         [ ]
    'Total'       [ 8.0178]   [ 17]      [ ]         [ ]         [ ]
stats =
    gnames: {6x1 cell}
    n: [3 3 3 3 1 5]
    source: 'anova1'
    means: [13.3333 14.2000 13.1000 13.4667 13.4000 14.1400]
    df: 12
    s: 0.6168
```

ANOVA Table

| Source | SS | df | MS | F | Prob>F |
|--------|---------|----|---------|------|--------|
| ----- | | | | | |
| Groups | 3.45244 | 5 | 0.69049 | 1.81 | 0.1843 |
| Error | 4.56533 | 12 | 0.38044 | | |
| Total | 8.01778 | 17 | | | |

Click on the group you want to test



-
1. The treatment populations must be normal.
 2. The treatment populations must all have the same variance

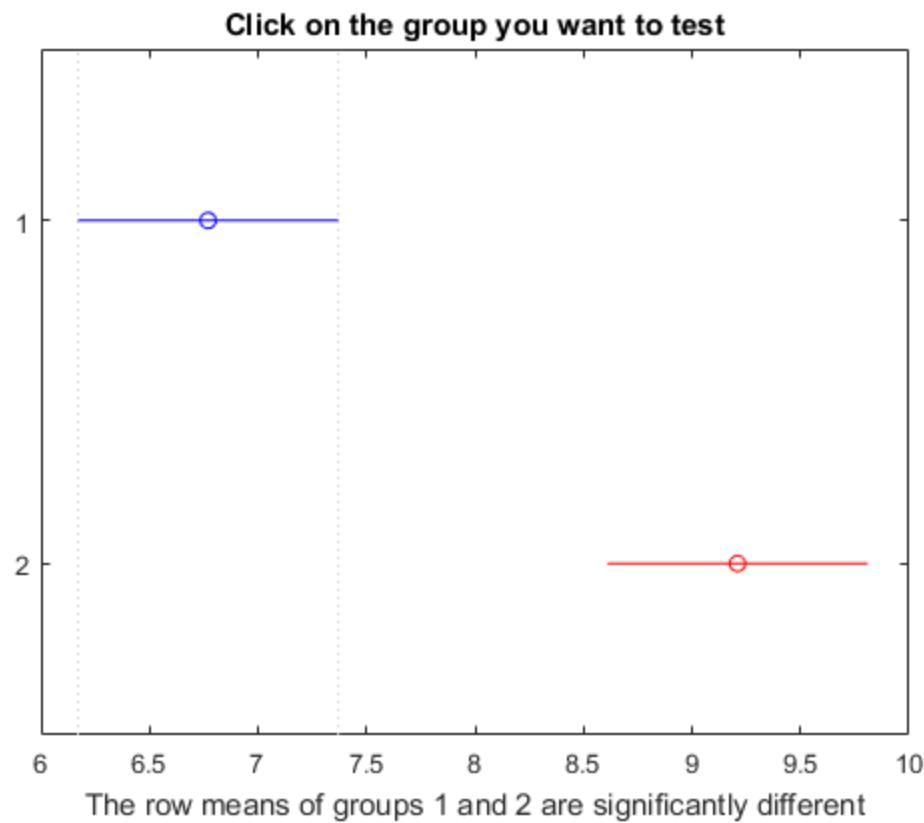
9.3-6

```
SA =[6.4 5.8 5.1 8.4 7.0 8.4 8.5 7.5 7.0 7.9]';
SB =[4.7 4.7 3.8 5.3 10.6 4.5 8.2 10.8 5.1 5.7]';
IA =[11.0 8.9 9.3 9.2 7.9 9.7 9.0 12.5 6.7 9.8]';
IB =[8.9 7.0 10.7 10.3 6.2 12.2 7.0 9.5 8.7 9.7]';

data = [SA SB; IA IB];
[p_vals, table,stats] = anova2(data, 10)
multcompare(stats,'estimate','column');
multcompare(stats,'estimate','row');

% Yes, it is possible, P=0.687
% Yes, there's a difference, p=0.00002
% Yes, no difference, p=0.3014

p_vals =
    0.3014    0.0002    0.6873
table =
    Columns 1 through 5
    'Source'      'SS'        'df'        'MS'        'F'
    'Columns'     [ 3.8440]   [ 1]     [ 3.8440]   [ 1.0992]
    'Rows'        [ 59.5360]  [ 1]     [59.5360]  [17.0238]
    'Interaction' [ 0.5760]   [ 1]     [ 0.5760]   [ 0.1647]
    'Error'       [125.9000]  [36]     [ 3.4972]   []
    'Total'       [189.8560]  [39]           []           []
    Column 6
    'Prob>F'
    [ 0.3014]
    [2.0838e-04]
    [ 0.6873]
    []
    []
stats =
    source: 'anova2'
    sigmasq: 3.4972
    colmeans: [8.3000 7.6800]
    coln: 20
    rowmeans: [6.7700 9.2100]
   rown: 20
    inter: 1
    pval: 0.6873
    df: 36
Note: Your model includes an interaction term. A test of main effects
can be
difficult to interpret when the model includes interactions.
Note: Your model includes an interaction term. A test of main effects
can be
difficult to interpret when the model includes interactions.
```



9.3-9

```

S80T5 =[5 6 5 5 4 3]';
S80T10 =[8 8 8 8 8 8]';
S80T15 =[11 10 9 9 10 9]';
S150T5 =[9 11 9 8 10 9]';
S150T10 =[14 14 15 13 17 18]';
S150T15 =[16 15 26 24 24 25]';
S220T5 =[34 33 19 21 18 20]';
S220T10 =[60 59 29 31 28 31]';
S220T15 =[65 64 31 33 75 80]';

data = [S80T5 S80T10 S80T15;S150T5 S150T10 S150T15;S220T5 S220T10
S220T15];
[p_vals, table,stats] = anova2(data, 6);

% No, it is not possible, P=0.007
% No, no model
% No, no model

```

| ANOVA Table | | | | | |
|-------------|---------|----|---------|-------|--------|
| Source | SS | df | MS | F | Prob>F |
| <hr/> | | | | | |
| Columns | 2619.1 | 2 | 1309.57 | 15.44 | 0 |
| Rows | 10795.8 | 2 | 5397.91 | 63.65 | 0 |
| Interaction | 1357.5 | 4 | 339.38 | 4 | 0.0073 |
| Error | 3816.3 | 45 | 84.81 | | |
| Total | 18588.8 | 53 | | | |

Because the design is not balanced, the analysis becomes harder, and we do not know how to do the calculations.

9.5-4

```

data = [0.55, 0.49...
        0.60, 0.42...
        0.37, 0.28...
        0.30, 0.28...
        0.54, 0.54...
        0.54, 0.47...
        0.44, 0.33...
        0.36, 0.20];

f=ff2n(3+1);
A=f(:,3)';
B=f(:,2)';
C=f(:,1)';

[p,table] = anovan(data,{A,B,C}, 'model','full', 'varnames',...
{'A','B','C'})

% Yes, it is appropriate (no interactions)
% B has an effect

p =

```

```

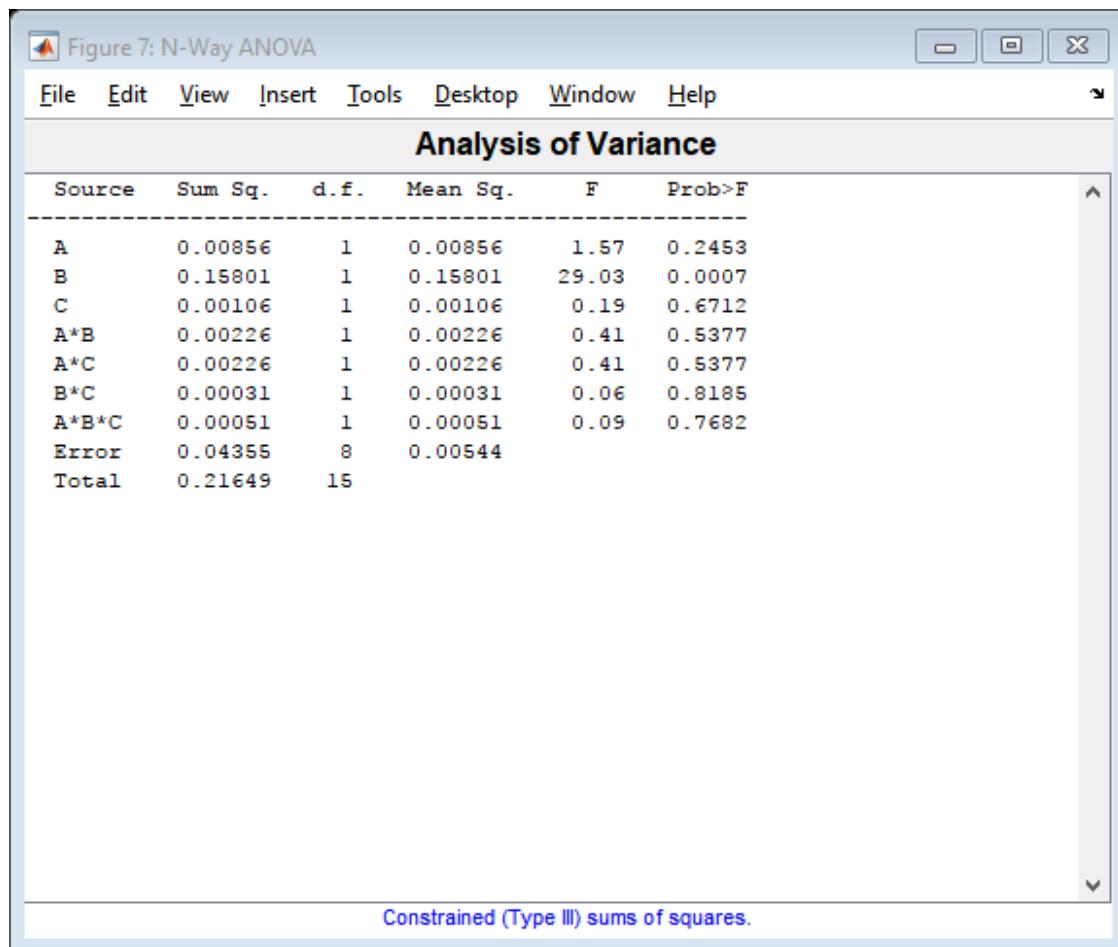
0.2453
0.0007
0.6712
0.5377
0.5377
0.8185
0.7682


| Columns 1 through 5 |               |        |             |               |
|---------------------|---------------|--------|-------------|---------------|
| 'Source'            | 'Sum Sq.'     | 'd.f.' | 'Singular?' | 'Mean Sq.'    |
| 'A'                 | [ 0.0086]     | [ 1]   | [ 0]        | [ 0.0086]     |
| 'B'                 | [ 0.1580]     | [ 1]   | [ 0]        | [ 0.1580]     |
| 'C'                 | [ 0.0011]     | [ 1]   | [ 0]        | [ 0.0011]     |
| 'A*B'               | [ 0.0023]     | [ 1]   | [ 0]        | [ 0.0023]     |
| 'A*C'               | [ 0.0023]     | [ 1]   | [ 0]        | [ 0.0023]     |
| 'B*C'               | [ 3.0625e-04] | [ 1]   | [ 0]        | [ 3.0625e-04] |
| 'A*B*C'             | [ 5.0625e-04] | [ 1]   | [ 0]        | [ 5.0625e-04] |
| 'Error'             | [ 0.0436]     | [ 8]   | [ 0]        | [ 0.0054]     |
| 'Total'             | [ 0.2165]     | [ 15]  | [ 0]        |               |



| 'F'        | 'Prob>F'      |
|------------|---------------|
| [ 1.5718]  | [ 0.2453]     |
| [ 29.0253] | [ 6.5572e-04] |
| [ 0.1940]  | [ 0.6712]     |
| [ 0.4145]  | [ 0.5377]     |
| [ 0.4145]  | [ 0.5377]     |
| [ 0.0563]  | [ 0.8185]     |
| [ 0.0930]  | [ 0.7682]     |
| [ ]        | [ ]           |
| [ ]        | [ ]           |


```



9.5-5

```

data = [68.0 77.5 98.0 98.0 74.0 77.0 97.0 98.0];
f = fliplr(ff2n(3));
f(f==0)=-1;
f(:,4)=f(:,1).*f(:,2);
f(:,5)=f(:,1).*f(:,3);
f(:,6)=f(:,2).*f(:,3);
f(:,7)=f(:,1).*f(:,2).*f(:,3);

effect = data*f/4

% No, design is unreplicated
% No, B has the greatest effect

effect =
    3.3750    23.6250    1.1250   -2.8750   -1.3750   -1.6250    1.8750

```

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