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%Name: David George
%StudentID:251004839
T = input_data();
%To conduct anova testing,the population must be normally distributed,
%independenat, and there most be homoegenity of vareinces between
groups.

%Checking to see if the population is normally distributed on the 12th
and 21st day.
%Kolmogorov Smirnov Test:
% H0: "the data is normally distributed"

% Rescale data for Norm(0,1) comparison 12th day:

    day12TestD1 = (( T.weight(T.Time ==12 & T.Diet ==
1)) - mean(T.weight(T.Time ==12 & T.Diet == 1))) /
(sqrt(var(T.weight(T.Time ==12 & T.Diet ==1))));
    day12TestD2 = (( T.weight(T.Time ==12 & T.Diet ==
2)) - mean(T.weight(T.Time ==12 & T.Diet == 2))) /
(sqrt(var(T.weight(T.Time ==12 & T.Diet ==2))));
    day12TestD3 = (( T.weight(T.Time ==12 & T.Diet ==
3)) - mean(T.weight(T.Time ==12 & T.Diet == 3))) /
(sqrt(var(T.weight(T.Time ==12 & T.Diet ==3))));
    day12TestD4 = (( T.weight(T.Time ==12 & T.Diet ==
4)) - mean(T.weight(T.Time ==12 & T.Diet == 4))) /
(sqrt(var(T.weight(T.Time ==12 & T.Diet ==4))));

% H: 0 (cannot reject) or 1 (reject: data unlikely normal)

[rejectday12D1 p12D1] = kstest(day12TestD1, 'Alpha', 0.05);
[rejectday12D2 p12D2] = kstest(day12TestD2, 'Alpha', 0.05);
[rejectday12D3 p12D3] = kstest(day12TestD3, 'Alpha', 0.05);
[rejectday12D4 p12D4] = kstest(day12TestD4, 'Alpha', 0.05);

    if rejectday12D1 == 0 & rejectday12D2 == 0 & rejectday12D3 == 0 &
rejectday12D4 == 0

        disp("Population on the 12th day is normally distriubuted");
    end

% Rescale data for Norm(0,1) comparison 21th day:

    day21TestD1 = (( T.weight(T.Time ==21 & T.Diet ==
1)) - mean(T.weight(T.Time ==21 & T.Diet == 1))) /
(sqrt(var(T.weight(T.Time ==21 & T.Diet ==1))));

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    day21TestD2 = (( T.weight(T.Time ==21 & T.Diet ==
2)) - mean(T.weight(T.Time ==21 & T.Diet == 2))) /
(sqrt(var(T.weight(T.Time ==21 & T.Diet ==2))));
    day21TestD3 = (( T.weight(T.Time ==21 & T.Diet ==
3)) - mean(T.weight(T.Time ==21 & T.Diet == 3))) /
(sqrt(var(T.weight(T.Time ==21 & T.Diet ==3))));
    day21TestD4 = (( T.weight(T.Time ==21 & T.Diet ==
4)) - mean(T.weight(T.Time ==21 & T.Diet == 4))) /
(sqrt(var(T.weight(T.Time ==21 & T.Diet ==4))));

[rejectday21D1 p21D1] = kstest(day21TestD1, 'Alpha', 0.05);
[rejectday21D2 p21D2] = kstest(day21TestD2, 'Alpha', 0.05);
[rejectday21D3 p21D3] = kstest(day21TestD3, 'Alpha', 0.05);
[rejectday21D4 p21D4] = kstest(day21TestD4, 'Alpha', 0.05);

% H: 0 (cannot reject) or 1 (reject: data unlikely normal)

    if rejectday21D1 == 0 & rejectday21D2 == 0 & rejectday21D3 == 0 &
rejectday21D4 == 0

        disp("Population on the 21st day is normally distriubuted");
    end

%Checking to determine homongenitity of variences between  diets on
day 12
%and 21 using the levene test

p12Var = vartestn(T.weight(T.Time ==12), T.Diet(T.Time
==12), 'TestType', 'LeveneAbsolute', 'Display', 'off');
p21Var = vartestn(T.weight(T.Time ==21), T.Diet(T.Time
==21), 'TestType', 'LeveneAbsolute', 'Display', 'off');

if p12Var > 0.05

    disp("The groups of chicks on the 12th, divided by diets, have
homogoenous variences");
end

if p21Var > 0.05

    disp("The groups of chicks on the 21st day, divided by diets,
have homogoenous variences");
end

% For both day 12 and day 21, the independence assumption holds up
because
% the probaility of one event occuring in no away affects the
probaility of

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% another event. In this case, the weight of any given chick in any
% group,
% has no bearing on the weight of any given chick in another group,
% nor
% does knowing the weight of one chick give insight into the probability
% of
% any other chick being a certain weight either. Therefore the
% independence
% assumption holds up on both the 12th and 21st day.

% Since all assumptions (normality, independence, and homogeneity of
% variances) are held up, anova testing can be performed, with post-hoc
% testing to determine the best diet.

% Anova testing and Post Hoc testing for day 12:

[pval12 tab12 stats12] = anova1(T.weight(T.Time ==12),
T.Diet(T.Time ==12), 'off');

figure

[c12,m12,h12,nms12] = multcompare(stats12, 'Alpha',
0.05, 'CType','bonferroni');
title("Day 12");
xlabel("Weight (grams)");
ylabel("Diet Number");

% Anova testing and Post Hoc testing for day 21:

[pval21 tab21 stats21] = anova1(T.weight(T.Time ==21),
T.Diet(T.Time ==21), 'off');

figure

[c21,m21,h21,nms21] = multcompare(stats21, 'Alpha',
0.05, 'CType','bonferroni');
title("Day 21");
xlabel("Weight (grams)");
ylabel("Diet Number");

Day12statarray = grpstats(T(T.Time
==12, :), 'Diet', 'mean', 'DataVars', {'weight'});
Day21statarray = grpstats(T(T.Time
==21, :), 'Diet', 'mean', 'DataVars', {'weight'});

Day12statarray.Properties.VariableNames{'Diet'} = 'DietsForDay12';
Day12statarray.Properties.VariableNames{'GroupCount'}
= 'GroupCountDay12';
Day12statarray.Properties.VariableNames{'mean_weight'}
= 'mean_weight_day12';

Day21statarray.Properties.VariableNames{'Diet'} = 'DietsForDay21';

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    Day21statarray.Properties.VariableNames{'GroupCount'}
= 'GroupCountDay21';
    Day21statarray.Properties.VariableNames{'mean_weight'}
= 'mean_weight_day21';
    both = [ Day12statarray Day21statarray ];
    disp(both);

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## Recomondations:

If the farmer is to select a chick on day 12, I advice the farmer to choose either diet 3 or diet 4. Looking at the figure for day 12, diet 3 and diet 4 are the only diets with statiscially differnt mean weights from diet 1, so either or works.

%If the farmer is to select a chick on day 21, I advice the farmer to  
 %choose diet 3. Looking at the figure for day 21, diet 3 is the only  
 diet  
 %that is statistically differnt from diet one.

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function T = input_data()
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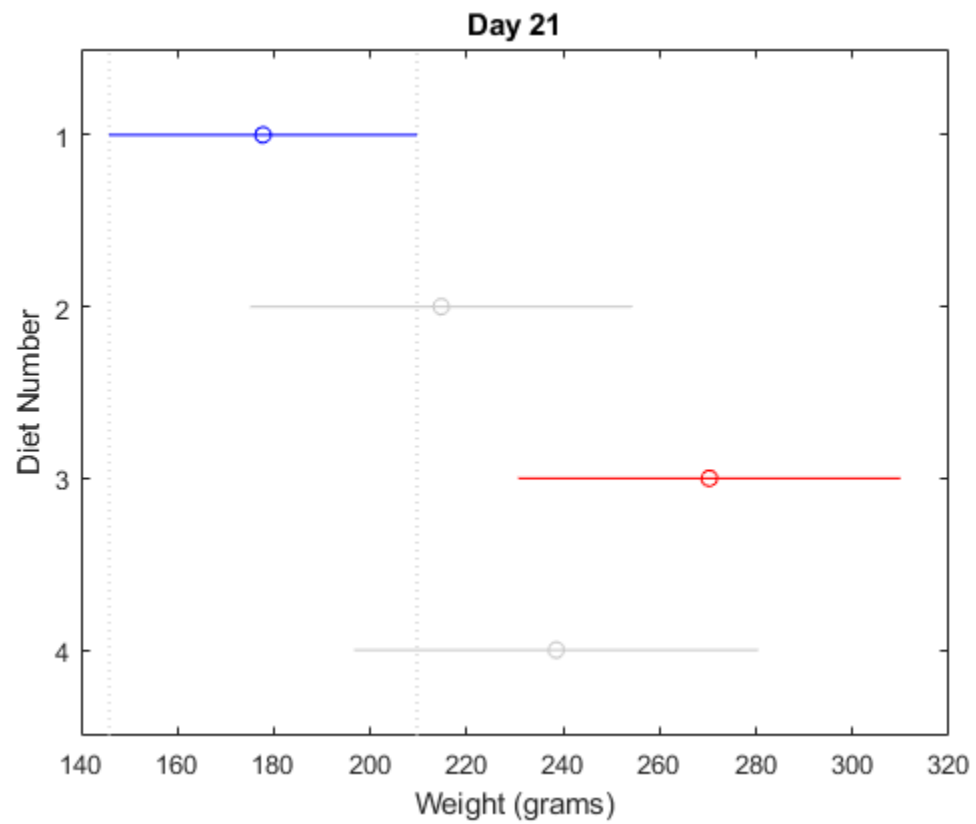
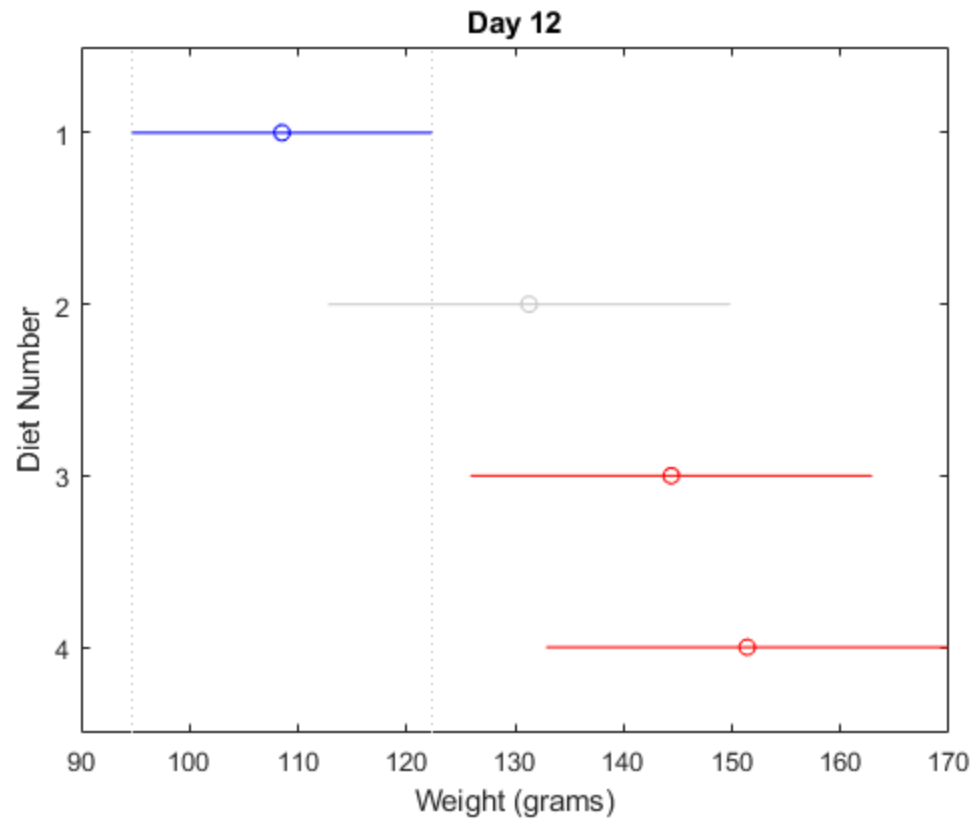
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%This function reads the input file and casts the data properly
T = readtable("chicks.csv");
T.weight = double(T.weight);
T.Time = double(T.Time);
T.Chick = uint32(T.Chick);
T.Diet = uint32(T.Diet);
end

```

*Population on the 12th day is normally distriubuted*  
*Population on the 21st day is normally distriubuted*  
*The groups of chicks on the 12th, divided by diets, have homogenous*  
*variances*  
*The groups of chicks on the 21st day, divided by diets, have*  
*homogenous variances*

	<i>DietsForDay12</i>	<i>GroupCountDay12</i>	<i>mean_weight_day12</i>
<i>DietsForDay21</i>	<i>GroupCountDay21</i>	<i>mean_weight_day21</i>	
1	1	19	108.526315789474
1	16		177.75
2	2	10	131.3
2	10		214.7
3	3	10	144.4
3	10		270.3
4	4	10	151.4
4	9		238.555555555556



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