```
Author: Jonathan W. Duggins
Last Update: 2016-01-27
Notes: Create ROOT data set for Lab 4
options nodate pageno=1 formdlim="~";
data root;
  input pretreat replicate variety length;
  cards;
    1 1
          11
  1
     1 2
  1
          26
  1
    1 3
          17
  1
     1 4 08
    2 1
  1
          05
  1
     2 2
          13
    2
  1
       3
          30
  1
    2 4
          05
  1
     3
       1
          07
    3 2
  1
          15
     3
       3
  1
          21
  1
     3
       4
          05
  2
     1 1
          15
  2
     1 2
          20
  2
     1 3
          15
  2
     1 4
          15
     2 1
          17
  2
     2 2
          21
  2
     2
       3
          29
  2
    2 4
         12
  2
    3
       1
          04
  2
    3 2
          20
  2
     3 3
          28
  2
     3 4
          10
  3
    1 1 03
  3
     1 2
          05
  3
     1 3
          06
  3
     1 4
          10
  3
     2 1
          01
  3
     2 2
         04
  3
     2
       3
          03
  3
    2 4
          10
  3
     3
       1
          06
  3
     3
       2
         04
  3
     3 3
          04
  3
     3 4
          05
run;
```

/****************

```
/*Put your code here!*/
/*Two-way ANOVA model*/
proc glm data = root plots=all ;
   class pretreat variety;
   model length = pretreat|variety/ alpha=0.00833 clparm;
   * benfoerroni mc , alpha = .05/6 = 0.00833;
   * simple effects with tukey corrections;
   lsmeans pretreat*variety / adjust = tukey cl pdiff alpha=0.05;
   estimate 'mu11-mu12' intercept 0 pretreat 0 0 0 variety 1 -1 0
pretreat*variety 1 -1 0 0 0 0 0 0 0 0 0
   estimate 'mu12-mu22' intercept 0 pretreat 1 -1 0 variety 0 0 0
pretreat*variety 0 1 0 0 0 -1 0 0 0 0 0
   estimate 'mu14-mu24' intercept 0 pretreat 1 -1 0 variety 0 0
pretreat*variety 0 0 0 1
                          0 0 0 -1 0 0 0 0 ;
   estimate 'mu13-mu23' intercept 0 pretreat 1 -1 0 variety 0 0 0
pretreat*variety 0 0 1 0 0 0 -1 0 0 0 0 0 ;
   * pretreat has three categories so we ahve three splits for alpha,
variety has four types so we have four betas;
run;
quit;
proc glm data = root plots = none;
   class pretreat variety;
   model length = pretreat|variety;
   * what is the effect on each pretrat of variety;
   lsmeans pretreat*variety / cl slice = pretreat;
   * what is the effect on each variety of pretreat;
   lsmeans pretreat*variety / slice = variety;
run;
quit;
```