How to calculate the mean

KE Lotterhos
3/9/2021

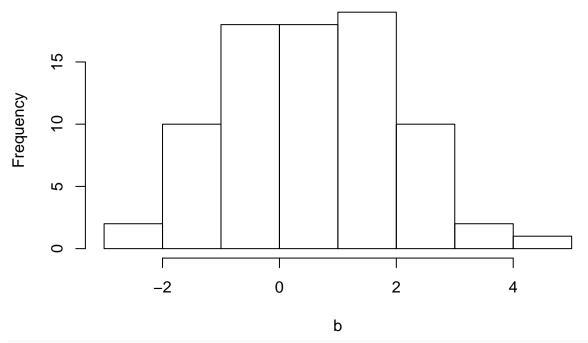
```
require(lme4)
## Loading required package: lme4
## Loading required package: Matrix
require(emmeans)
## Loading required package: emmeans
## Welcome to emmeans.
## NOTE -- Important change from versions <= 1.41:
##
       Indicator predictors are now treated as 2-level factors by default.
       To revert to old behavior, use emm_options(cov.keep = character(0))
# Different ways of calculating means ####
getmeans <- function(ind=1:length(a), true){</pre>
  # model fit
  G <- as.factor(G_lev[ind])</pre>
  E <- as.factor(E_lev[ind])</pre>
  m1 \leftarrow lm(b[ind] \sim G * E)
  emms <- as.data.frame(emmeans(m1, ~ G * E))
  emmG <- as.data.frame(emmeans(m1, "G"))</pre>
  emmE <- as.data.frame(emmeans(m1, "E"))</pre>
  obs_all <- mean(b[ind]) # This may or may not be what your eMM does
  obs_mean_GE <- mean(tapply(b[ind],a[ind], mean))</pre>
  obs_mean_G <- mean(tapply(b[ind],G_lev[ind], mean))</pre>
  obs_mean_E <- mean(tapply(b[ind],E_lev[ind], mean))</pre>
  obs_mean_cGE <- mean(c(obs_mean_G, obs_mean_E)) # Note they count unequally in the common garden N x
  EMM_mean_GE <- mean(emms$emmean)</pre>
  EMM_mean_G <- mean(emmG$emmean)</pre>
  EMM_mean_E <- mean(emmE$emmean)</pre>
  EMM_mean_cGE <- mean(c(EMM_mean_E,EMM_mean_G))</pre>
  out <- cbind(obs_all, obs_mean_GE, obs_mean_E, obs_mean_cGE, EMM_mean_GE, EMM_mean_G, EMM_
  round(out - true, 4)
```

consider 2x2 case

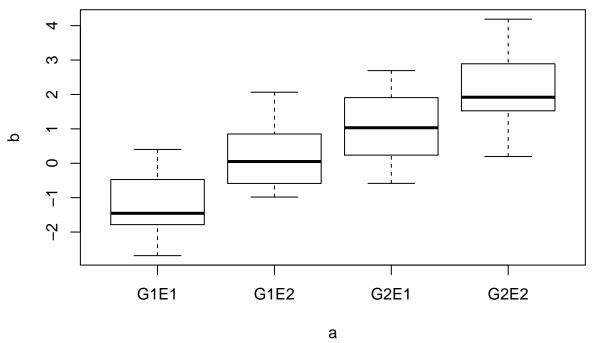
First, consider the 2x2 case, with different GE means for each level. Compare results for imbalance in the level with the lowest mean, and imbalance in the level with the highest mean.

```
a <- rep(c("G1E1","G1E2", "G2E1", "G2E2"), each=20)
G_lev <- substr(a, start=1, stop=2)
E_lev <- substr(a, start=3, stop=4)
b <- c(rnorm(length(a)/4, -1), rnorm(length(a)/4, 0), rnorm(length(a)/4, 1), rnorm(length(a)/4, 2))
hist(b)</pre>
```

Histogram of b



boxplot(b~a)



```
true1 <- mean(c(-1,0,1,2)) # true mean
getmeans(1:length(a), true1) # no imbalance</pre>
```

NOTE: Results may be misleading due to involvement in interactions
NOTE: Results may be misleading due to involvement in interactions
obs_all obs_mean_GE obs_mean_E obs_mean_cGE EMM_mean_GE

```
## [1,] 0.0476
                     0.0476
                                0.0476
                                         0.0476
                                                      0.0476
                                                                    0.0476
       EMM_mean_G EMM_mean_E EMM_mean_cGE
                      0.0476
            0.0476
getmeans(15:length(a), true1) # imbalance in first category
## NOTE: Results may be misleading due to involvement in interactions
## NOTE: Results may be misleading due to involvement in interactions
        obs all obs mean GE obs mean G obs mean E obs mean cGE EMM mean GE
                                           0.3394
                                                        0.2818
## [1,] 0.4053
                     0.0268
                                0.2242
##
        EMM mean G EMM mean E EMM mean cGE
## [1,]
            0.0268
                      0.0268
                                    0.0268
getmeans(1:(length(a)-15), true1) # imbalance in last category
## NOTE: Results may be misleading due to involvement in interactions
## NOTE: Results may be misleading due to involvement in interactions
##
        obs_all obs_mean_GE obs_mean_G obs_mean_E obs_mean_cGE EMM_mean_GE
## [1,] -0.2715
                     0.168
                               -0.0582
                                          -0.1865
                                                       -0.1223
        EMM_mean_G EMM_mean_E EMM_mean_cGE
##
             0.168
                        0.168
## [1,]
                                     0.168
par(mar=c(10,3,1,1))
bob <- replicate(100,getmeans(c(sample(1:20,5),20:length(a)), true1),simplify = TRUE)
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rownames(bob) <- c("obs_all", "obs_mean_GE", "obs_mean_G", "obs_mean_E", "obs_mean_cGE", "EMM_mean_GE",
#head(bob)
boxplot(t(bob), las=2)
abline(h=0)
 0.4
 0.3
 0.2
```

bob <- replicate(100,getmeans(c(1:(length(a)-20), sample((length(a)-20):length(a),5)), true1),simplify

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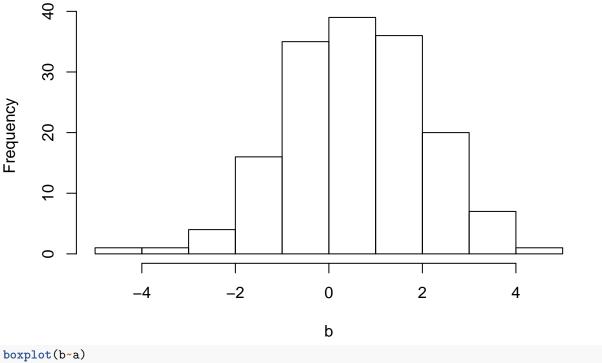
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#head(bob)
rownames(bob) <- c("obs_all", "obs_mean_GE", "obs_mean_G", "obs_mean_E", "obs_mean_cGE", "EMM_mean_GE",
#head(bob)
boxplot(t(bob), las=2)
abline(h=0)
 0.2
 0.1
 0.0
-0.1
-0.2
-0.3
-0.4
                                       obs_mean_E
                      obs_mean_GE
                                               obs_mean_cGE
                                                        EMM_mean_GE
                                                                                 EMM_mean_cGE
                               obs_mean_
                                                                EMM_mean_
                                                                         EMM mean
```

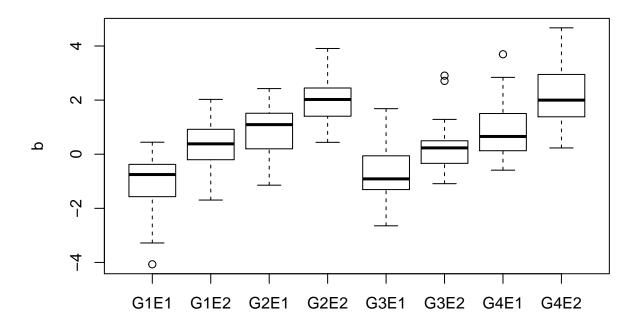
consider 2x4 common garden case

First, consider the 2x4 case, with different GE means for each level. Compare results for imbalance in the level with the lowest mean, and imbalance in the level with the highest mean.

```
a <- rep(c("G1E1", "G1E2", "G2E1", "G2E2", "G3E1", "G3E2", "G4E1", "G4E2"), each=20)
G_lev <- substr(a, start=1, stop=2)</pre>
E_lev <- substr(a, start=3, stop=4)</pre>
b <- c(rnorm(length(a)/8, -1),
       rnorm(length(a)/8, 0),
       rnorm(length(a)/8, 1),
       rnorm(length(a)/8, 2),
       rnorm(length(a)/8, -1),
       rnorm(length(a)/8, 0),
       rnorm(length(a)/8, 1),
       rnorm(length(a)/8, 2))
hist(b)
```

Histogram of b





а

```
## NOTE: Results may be misleading due to involvement in interactions
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rownames(bob) <- c("obs_all", "obs_mean_GE", "obs_mean_G", "obs_mean_E", "obs_mean_cGE", "EMM_mean_GE",
#head(bob)
boxplot(t(bob), las=2)
abline(h=0)
0.25
0.20
0.15
0.10
0.05
0.00
                     obs_mean_GE
                                      s_mean_
bob <- replicate(100,getmeans(c(1:(length(a)-20),
                                sample((length(a)-20):length(a),5)), true2),
                 simplify = TRUE)
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