Leveraging multi-harvest data for simultaneously selecting high-yield and witches' broom-resistant Theobroma grandiflorum hybrids

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Required packages	

Required packages

```
require(asreml)
require(tidyverse)
require(patchwork)
```

Loading the data

```
data = read.csv("data.csv", header = T, sep=';')
```

Setting the factors

Linear Mixed Models

Homoscedastic model for Fruit Yield

	Value	Std. Error	Conf. Int.
σ_q^2	14.45	5.56	10.90
$\sigma_{g}^{2} \ \sigma_{p}^{2} \ \sigma_{gh}^{2} \ \sigma_{e}^{2}$	35.08	4.24	8.31
σ_{qh}^2	12.02	1.99	3.90
σ_e^2	71.08	2.47	4.84
r	0.82	NA	NA
H_g^2	0.68	NA	NA

Heteroscedastic model for Fruit Yield

```
fy2 = asreml(fixed = yd ~ Harvests + Replicates:Harvests,
             random = ~ Hybrids + Hybrids:Harvests + Plots,
             residual = ~dsum(~id(units)|Harvests),
             data = data)
sum.fy2 = summary(fy2) varcomp[,1:2]
sum.fy2$CI = sum.fy2$std.error * 1.96
aic.fy2 = summary(fy2)$aic
predfy2_vcov = predict(fy2, classify = "Hybrids", vcov = T)
predfy2_sed = predict(fy2, classify = "Hybrids", sed = T)
PEV = mean(diag(predfy2_vcov$vcov))
MVdelta = mean((predfy2_sed$sed^2)[upper.tri(predfy2_sed$sed^2,diag = F)])
acc2 = sqrt(1-(PEV/sum.fy2[1,1]))
her2 = 1-(MVdelta/(2*sum.fy2[1,1]))
# Harvest-wise heritability
her2j = NULL
for(i in 1:nlevels(data$Harvests)){
 predfy2_sed = predict(fy2, classify = "Hybrids:Harvests",
                        level=list(Harvests = i), sed = T)
 MVdelta = mean((predfy2_sed$sed^2)[upper.tri(predfy2_sed$sed^2,diag = F)])
 her2j[i] = 1-(MVdelta/(2*sum.fy2[1,1]))
}
blups.fy = coef(fy2)$random[grep("Hybrids", rownames(coef(fy2)$random))]
```

	Value	Std. Error	Conf. Int.
σ_a^2	13.88	4.93	9.67
$\sigma_n^{\frac{5}{2}}$	26.62	3.23	6.33
σ_{ab}^2	7.92	1.56	3.05
$\sigma_{e_1}^2$	28.98	3.43	6.72
$\sigma_{e_2}^{ ilde{2}^1}$	28.41	3.36	6.59
$\sigma_{e_3}^2$	40.46	4.43	8.69
$\sigma_{e_{A}}^{ ilde{2}^{3}}$	49.32	5.27	10.33
$\sigma_{e_5}^{ ilde{2}^{st}}$	72.59	7.46	14.62
$\sigma_{e_6}^{ ilde{2}^\circ}$	98.50	10.09	19.79
$\sigma_{e_7}^{ ilde{2}^0}$	167.26	16.52	32.38
$\sigma_{e_8}^2$	101.13	10.43	20.44
$\sigma_{e_0}^{ ilde{2}^\circ}$	100.17	10.05	19.69
r	0.85	0.85	0.85
$H_{a_1}^2$	0.57	0.57	0.57
$H_{g_2}^{\stackrel{3}{2}}$	0.57	0.57	0.57
$H_{a_2}^2$	0.52	0.52	0.52
$H_{a_4}^{\widetilde{23}}$	0.49	0.49	0.49
$H_{a_5}^{2^*}$	0.43	0.43	0.43
$H_{q_e}^{\widetilde{2}\widetilde{\circ}}$	0.39	0.39	0.39
$egin{array}{l} \sigma_{g}^{2} \ \sigma_{gh}^{2} \ \sigma_{gh}^{2} \ \sigma_{e_{1}}^{2} \ \sigma_{e_{2}}^{2} \ \sigma_{e_{3}}^{2} \ \sigma_{e_{5}}^{2} \ \sigma_{e_{6}}^{2} \ \sigma_{e_{7}}^{2} \ \sigma_{e_{9}}^{2} \ r \ H_{g_{1}}^{2} \ H_{g_{2}}^{2} \ H_{g_{5}}^{2} \ H_{g_{6}}^{2} \ H_{g_{7}}^{2} \ H_{g_{9}}^{2} \ H_$	0.31	0.31	0.31
H_{as}^2	0.38	0.38	0.38
$H_{q_0}^{2^{\circ}}$	0.38	0.38	0.38

Coefficients of determination

```
sig2f = NULL
cgh = NULL
cp = NULL

for (i in 1:nlevels(data$Harvests)) {
    sig2f[i] = sum(sum.fy2[1:3,1]) + sum.fy2[i+3,1]
    cgh[i] = sum.fy2["Hybrids:Harvests", "component"] / sig2f[i]
    cp[i] = sum.fy2["Plots", "component"] / sig2f[i]
    ce[i] = sum.fy2[i+3,1] / sig2f[i]
}

coef.det = data.frame(
    "Harvests" = levels(data$Harvests),
    "sig2f" = sig2f,
    "chg" = cgh,
    "cp" = cp,
```

```
"ce" = ce
)
```

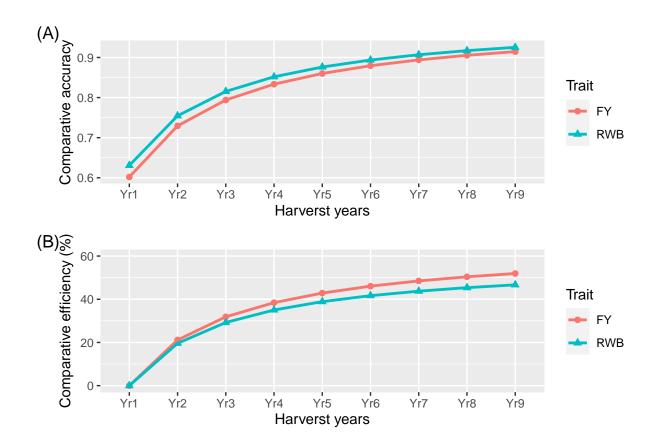
Harvests	σ_f^2	c_{gh}^2	c_p^2	c_e^2
Yr1	77.40	0.10	0.34	0.37
Yr2	76.83	0.10	0.35	0.37
Yr3	88.88	0.09	0.30	0.46
Yr4	97.74	0.08	0.27	0.50
Yr5	121.01	0.07	0.22	0.60
Yr6	146.92	0.05	0.18	0.67
Yr7	215.68	0.04	0.12	0.78
Yr8	149.56	0.05	0.18	0.68
Yr9	148.59	0.05	0.18	0.67

Generalized linear mixed model for witches' broom resistance

Optimum number of harvests

```
rho_fy = NULL
for (i in 1:nlevels(data$Harvests)) {
   rho_fy[i] = sum(sum.fy2["Hybrids","component"],sum.fy2["Plots","component"])/
       sig2f[i]
}
rho_fy = mean(rho_fy) #Fruit yield repeatability
```

```
rho wb = sum(sum.wb["Hybrids","component"],sum.wb["Plots","component"])/
    sum(c(sum.wb[1:3,1],3.29)) #WB resistance repeatability
effh fy = NULL
acch_fy = NULL
effh_wb = NULL
acch wb = NULL
for (i in 1:nlevels(data$Harvests)) {
  effh_fy[i] = sqrt(i/(1+(i-1)*rho_fy))
  effh_wb[i] = sqrt(i/(1+(i-1)*rho_wb))
  acch_fy[i] = sqrt((i*rho_fy)/(i*rho_fy+1-rho_fy))
  acch_wb[i] = sqrt((i*rho_wb)/(i*rho_wb+1-rho_wb))
}
a = data.frame(
  "Harvests" = rep(levels(data$Harvests),2),
  "Trait" = rep(c("FY", "RWB"), each = nlevels(data$Harvests)),
  "Accuracy" = c(acch_fy,acch_wb)
) %>% ggplot()+
  geom_point(aes(x = Harvests, y = Accuracy, color = Trait, shape = Trait),
             size=2)+
  geom line(aes(x = Harvests, y = Accuracy, color = Trait, group = Trait),
  labs(y = "Comparative accuracy",x = "Harverst years",tag = "(A)")
b = data.frame(
  "Harvests" = rep(levels(data$Harvests),2),
  "Trait" = rep(c("FY", "RWB"), each = nlevels(data$Harvests)),
  "Efficiency" = c((effh_fy-1)*100, (effh_wb-1)*100)
) %>% ggplot()+
  geom point(aes(x = Harvests, y = Efficiency, color = Trait, shape = Trait),
             size=2)+
  geom_line(aes(x = Harvests, y = Efficiency, color = Trait, group = Trait),
            size=1) + ylim(0,60) +
  labs(y = "Comparative efficiency (%)", x = "Harverst years", tag = "(B)")
a/b
```



Selection index

Obtaining the yield persistence

```
blups.fy = left_join(data.frame(
    'Hybrids' = rep(levels(data$Hybrids), each = nlevels(data$Harvests)),
    'Harvests' = rep(levels(data$Harvests), nlevels(data$Hybrids)),
    'blup' = blups.fy[(nlevels(data$Hybrids)+1):length(blups.fy)]
), data.frame(
    'Hybrids' = levels(data$Hybrids),
    'blup' = blups.fy[1:nlevels(data$Hybrids)]
), by='Hybrids') %>% mutate(
    BLUP = blup.x + blup.y
) %>% select(Hybrids, Harvests, BLUP)

max.fy = blups.fy %>% group_by(Harvests) %>%
    summarise(max = max(BLUP))

num = NULL
```

```
for (i in levels(data$Hybrids)) {
   dttes = blups.fy %>% filter(Hybrids == i) %>%
     select(Hybrids, Harvests, BLUP)
   num[i] = 1/sum((dttes$BLUP - max.fy$max)^2)
}
den = sum(num)^2
Per.fy = num/den
Per.fy = rownames_to_column(as.data.frame(Per.fy),'Hybrids')
```

Additive index

Hybrids	PERS.FY	BLUP.FY	BLUP.RWB	AI
H125	2.55	6.94	-0.64	2.39
H165	0.51	5.54	-0.68	2.05
H143	0.09	1.72	-1.17	1.43
H117	0.26	4.50	-0.09	1.28
H137	0.04	-0.76	-1.73	1.23
H140	0.26	4.48	0.05	1.16
H167	0.08	1.68	-0.85	1.15
H124	0.05	-0.37	-1.30	0.98
H129	0.11	2.55	-0.17	0.83
H127	0.08	1.43	-0.43	0.74
H149	0.05	-0.19	-0.89	0.69
H135	0.04	-1.60	-1.31	0.65
H157	0.04	-0.50	-0.89	0.60
H131	0.05	-0.05	-0.50	0.40
H166	0.04	-0.71	-0.65	0.35
H118	0.03	-2.09	-0.94	0.22
H130	0.06	0.39	-0.08	0.17
H132	0.03	-2.18	-0.65	-0.05
H162	0.15	3.46	1.46	-0.28
H169	0.28	4.57	1.86	-0.32
H120	0.12	2.53	1.23	-0.34
H134	0.03	-3.36	-0.65	-0.36
H133	0.03	-3.39	-0.52	-0.48
H136	0.02	-4.03	-0.68	-0.51
H144	0.08	1.39	1.10	-0.54
H163	0.08	1.44	1.38	-0.76
H138	0.05	-0.33	1.10	-1.00
H152	0.04	-1.71	0.91	-1.21
H150	0.04	-1.16	1.30	-1.39
H121	0.04	-1.60	1.29	-1.50
H161	0.06	0.40	1.96	-1.51
H128	0.02	-6.38	-0.20	-1.54
H123	0.01	-7.96	-0.07	-2.08
H126	0.02	-4.67	1.45	-2.46