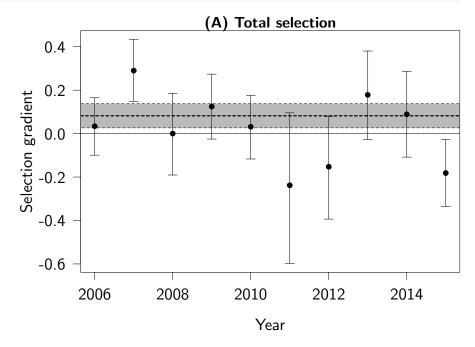
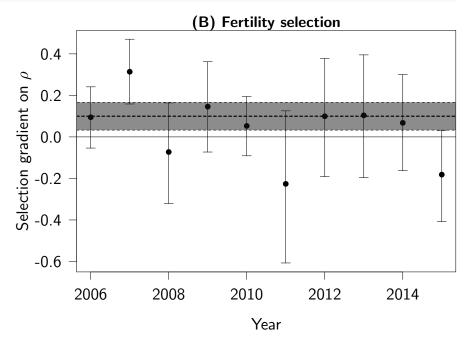
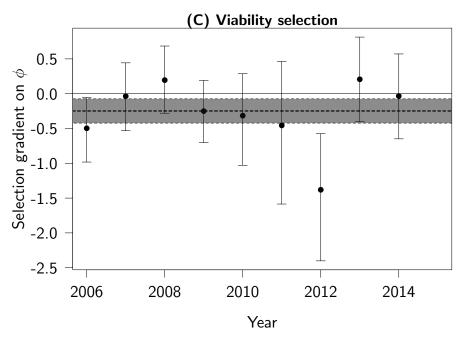
```
## Loading required package: Matrix
## Loading required package: coda
## Loading required package: ape
```



 $\#points (x=2006:2015, y=unlist (coefficients (mmRnoCorfitness) \pounds Year ["StMass"]), pch=17)$





Correlation fertility viability

```
cor.test(YearPheno$Phi,YearPheno$Rho)

##

## Pearson's product-moment correlation

##

## data: YearPheno$Phi and YearPheno$Rho

## t = -1.9473, df = 1292, p-value = 0.05171

## alternative hypothesis: true correlation is not equal to 0

## 95 percent confidence interval:

## -0.1082724891 0.0003989614

## sample estimates:

## cor

## -0.05409695
```

```
paste(round(smmRnoCorrho$coefficients[2,1],rounding)," (",round(smmRnoCorrho$coefficients[2,1])
             paste(round(smmRnoCorphi$coefficients[2,1],rounding)," (",round(smmRnoCorphi$coefficients[2,1])
SigmaA <- c(sqrt(as.numeric(smmARnoCorfitness$varcor$Year.1)),</pre>
             sqrt(as.numeric(smmRnoCorrho$varcor$Year.1)),
sqrt(as.numeric(smmRnoCorphi$varcor$Year.1)))
SigRat <- c(smmARnoCorfitness$coefficients[2,1]/sqrt(as.numeric(smmARnoCorfitness$varcor$Year
             smmRnoCorrho$coefficients[2,1]/sqrt(as.numeric(smmRnoCorrho$varcor$Year.1)),
             smmRnoCorphi$coefficients[2,1]/sqrt(as.numeric(smmRnoCorphi$varcor$Year.1)))
psigmaA <- c(fitnessAanova$`Pr(>Chisq)`[2]/2, RhoAanova$`Pr(>Chisq)`[2]/2, PhiAanova$`Pr(>Cl
confsigma <- c(paste("[",round(CImmARnoCorfitness[2,1],rounding),";",round(CImmARnoCorfitnes
                paste("[",round(CImmRnoCorrho[2,1],rounding),";",round(CImmRnoCorrho[2,2],round)
                paste("[",round(CImmRnoCorphi[2,1],rounding),";",round(CImmRnoCorphi[2,2],rounding))
TabSel <- data.frame(BetaGlm = BetaGlm, B=SDyears, C=SEyears , D=BetaGLMM , E=SigmaA, DD =co
                                Table 1:
 0.082 (0.028)
              0.167
                    0.097
                           0.036 (0.044)
                                        0.117
                                               [0.063;0.218]
                                                           8.1E-06
                                                                    0.309
 0.1 (0.034)
              0.160
                    0.117
                           0.052 (0.044)
                                        0.111
                                               [0.053;0.212]
                                                           2.5E-04
                                                                    0.466
 -0.248 (0.089)
              0.484
                    0.319
                           -0.217 (0.098)
                                        0.109
                                               [0;0.425]
                                                           3.6E-01
                                                                  -1.998
   Correlation between selection and evolution
posterior.mode(as.mcmc(SelToG))
##
       var1
## 0.360992
HPDinterval(as.mcmc(SelToG))
              lower
                        upper
## var1 -0.3910723 0.635243
## attr(,"Probability")
## [1] 0.95
szgr <- 2
```

par(las=1,mar=marr, cex=szgr, cex.lab=szax , cex.axis=szax, lwd=2 , las=1)

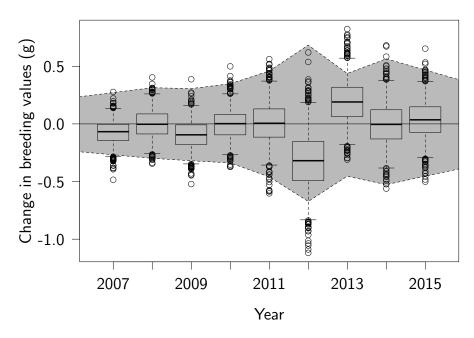
bbv <- boxplot(bvpairwise,ylab="Change in breeding values (g)", xlab="Year", range = 1,cex=:polygon(x = c(2006,2008:2014,2016,2016,2014:2008,2006) -2006, y = c(LowDrift,rev(HighDrit))

szax <- 1.3

bbv\$stats

marr $\leftarrow c(4, 4, 1, 1) + 0.1$

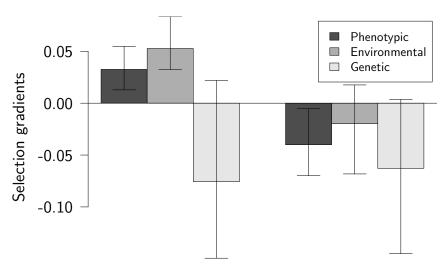
```
[,1] [,2] [,3] [,4] [,5]
## [1,] -0.280501568 -0.257383635 -0.347173588 -0.267832000 -0.357132486
## [2,] -0.143366410 -0.087176846 -0.178386364 -0.095028840 -0.114692786
## [3,] -0.067520536 -0.003132208 -0.094707610 0.002214962 0.005447066
## [5,] 0.134012211 0.260359089 0.161244353 0.260692898 0.371425991
##
     [,6]
          [,7]
                [,8]
## [1,] -0.8321504 -0.17794910 -0.382623528 -0.29096950
## [3,] -0.3192183  0.19033674 -0.004056911  0.03549705
## [4,] -0.1519085  0.31816724  0.122591670  0.14822783
## [5,] 0.1872504 0.57031481 0.374995899 0.36644027
bbv$group
##
  ##
  \hbox{ \#\# [246] 7777777777777777777777777777888888} 
## [351] 9
abline(h=0)
```



```
density(bvpairwise[,1])
##
## Call:
##
   density.default(x = bvpairwise[, 1])
##
## Data: bvpairwise[, 1] (1000 obs.); Bandwidth 'bw' = 0.02343
##
##
                             :0.000192
##
   Min.
         :-0.5547
                      Min.
##
   1st Qu.:-0.3292
                      1st Qu.:0.018199
##
   Median :-0.1036
                      Median :0.424183
##
   Mean
         :-0.1036
                      Mean
                             :1.107278
   3rd Qu.: 0.1219
                      3rd Qu.:2.109751
##
##
   Max. : 0.3475
                      Max. :3.900320
```

```
setPar()
par(mar=c(4, 6, 1, 1) + 0.1)
Betas <- matrix(sapply(X = list(BetaP1, BetaE1, BetaG1, BetaP2, BetaE2, BetaG2), posterior.n
BetasCI <- sapply(X = list(BetaP1, BetaE1, BetaG1, BetaP2, BetaE2, BetaG2), HPDinterval)

x <- barplot(Betas, beside=TRUE, ylim=c(min(BetasCI),max(BetasCI)),names.arg = c("Postive so abline(h=0)
arrows(x0 = x, y0=BetasCI[1,],y1=BetasCI[2,],angle = 90,code = 3)</pre>
```



```
posterior.mode(BetaG1 - BetaE1)
         var1
## -0.1234186
HPDinterval(BetaG1 - BetaE1)
##
             lower
                        upper
## var1 -0.2180723 -0.02770633
## attr(,"Probability")
## [1] 0.95
mean((BetaG1 - BetaE1)>0)*2
## [1] 0.006
posterior.mode(BetaG2 - BetaE2)
           var1
## -0.009610168
HPDinterval(BetaG2 - BetaE2)
```

```
## lower upper
## var1 -0.1379426 0.05755331
## attr(,"Probability")
## [1] 0.95
mean((BetaG2 - BetaE2)>0)*2
## [1] 0.424
posterior.mode(BetaE1 - BetaE2)
##
        var1
## 0.07575993
HPDinterval(BetaE1 - BetaE2)
            lower
                     upper
## var1 0.03828845 0.1372658
## attr(,"Probability")
## [1] 0.95
mean((BetaE1 - BetaE2)<0)*2</pre>
## [1] 0
posterior.mode(BetaG1 - BetaG2)
         var1
## -0.003689573
HPDinterval(BetaG1 - BetaG2)
##
             lower
                       upper
## var1 -0.08005378 0.07575714
## attr(,"Probability")
## [1] 0.95
mean((BetaG1 - BetaG2)>0)*2
## [1] 0.908
```

```
mat <- matrix(c(1, 2, 3, 4),ncol = 2, byrow = TRUE)
layout(mat, widths = rep.int(1, ncol(mat)), heights = c(1,1), respect = FALSE )
szgr <- 2
szax <- 1.3</pre>
```

```
par(las=1, cex=szgr, cex.lab=szax , cex.axis=szax, lwd=5 ,pch=16, las=1,
          oma=c(0,0,0,0)
barcol = grey(level = c(0.6,0.3))
nyears <- 6
h2 < -0.3
### Graph constant
par(mar=c(3, 4, 2, 1) + 0.1)
constant_selcov <- rep(1, times=nyears)</pre>
constant_evol <- constant_selcov * h2</pre>
constant_tp <- as.matrix(rbind(constant_selcov, constant_evol))</pre>
cumulative_constant <- cumsum(constant_evol)</pre>
xb1 <- barplot(height = constant_tp, ylim = c(-1.5,2), beside = TRUE,
                                      main="\\textbf{(A)}", yaxt="n", border = NA, col = barcol )
legend(x = "bottomleft", legend = c("Selection differential", "Genetic differential", "Evolution differential", "Evol
                                   bty = "o",
                  col = c(barcol, "black"),
                 lty = c(0, 0, 1), lwd = c(0, 0, 5),
                  pch = c(22, 22, NA),
                  pt.bg = c(barcol, NA),
                  pt.cex = 3, cex=1.5)
axis(side=2, at = c(-1,0,1,2))
axis(side=1, at = seq(from=min(xb1), to=max(xb1), length.out = nyears),
             labels = FALSE,tick = TRUE)
mtext(text = "Differential", side = 2, line = 2, cex=3, las=0)
abline(h=0)
lines(x=seq(from=min(xb1), to=max(xb1), length.out = nyears), y=cumulative_constant)
box(which = "plot", lty = "solid")
### Graph fluctuation
par(mar=c(3, 3, 2, 2) + 0.1)
flu_selcov \leftarrow c(1.2, 0.82, 0.6, 0.94, 1.3, 0.65)
flu_evol <- flu_selcov * h2
flu_tp <- as.matrix(rbind(flu_selcov, flu_evol))</pre>
cumulative_flu <- cumsum(flu_evol)</pre>
xb2 \leftarrow barplot(height = flu_tp, ylim = c(-1.5,2), beside = TRUE,
```

```
main="\\textbf{(B)}", yaxt="n", border = NA, col = barcol )
axis(side=2, at = c(-1,0,1,2), labels = FALSE, tick = TRUE)
axis(side=1, at = seq(from=min(xb1), to=max(xb1), length.out = nyears),
     labels = FALSE, tick = TRUE)
abline(h=0)
lines(x=seq(from=min(xb2), to=max(xb2), length.out = nyears), y=cumulative_flu)
box(which = "plot", lty = "solid")
### Graph reversal
par(mar=c(3, 4, 2, 1) + 0.1)
rev_selcov \leftarrow c(0.6,-1.1,-0.8,0.9,-0.22,0.5)
rev_evol <- rev_selcov * h2
rev_tp <- as.matrix(rbind(rev_selcov, rev_evol))</pre>
cumulative_rev <- cumsum(rev_evol)</pre>
xb3 <- barplot(height = rev_tp, ylim = c(-1.5,2), beside = TRUE,
               main="\\textbf{(C)}", yaxt="n", border = NA, col = barcol )
axis(side=2, at = c(-1,0,1,2))
axis(side=1, at = seq(from=min(xb1), to=max(xb1), length.out = nyears),
     labels = 1:6,tick = TRUE)
mtext(text = "Differential", side = 2, line = 2, cex=3, las=0)
abline(h=0)
lines(x=seq(from=min(xb3), to=max(xb3), length.out = nyears), y=cumulative_rev)
box(which = "plot", lty = "solid")
mtext(text = "Year", side = 1, line = 2, cex=3)
### Graph decoupling + flu
par(mar=c(3, 3, 2, 2) + 0.1)
dec_selcov \leftarrow c(0.6,-1.1,-0.8,0.9,-0.22,0.5)
dec_evol <- 0.3+dec_selcov * h2
dec_tp <- as.matrix(rbind(dec_selcov, dec_evol))</pre>
cumulative_dec <- cumsum(dec_evol)</pre>
xb5 \leftarrow barplot(height = dec_tp, ylim = c(-1.5,2), beside = TRUE,
               main="\\textbf{(D)}", yaxt="n", border = NA, col = barcol)
axis(side=2, at = c(-1,0,1,2), labels = FALSE, tick = TRUE)
axis(side=1, at = seq(from=min(xb1), to=max(xb1), length.out = nyears),
     labels = 1:6, tick = TRUE)
```

```
abline(h=0)
lines(x=seq(from=min(xb5), to=max(xb5), length.out = nyears), y=cumulative_dec)
box(which = "plot", lty = "solid")
mtext(text = "Year", side = 1, line = 2,cex = 3)
```

