

Supplementary material to: “Are animals shrinking due to climate change? Temperature-mediated selection on body mass in mountain wagtails”

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This document presents the code used for the analysis in: “Are animals shrinking due to climate change? Temperature-mediated selection on body mass in mountain wagtails”.

Trends in climate

We first explored the climate data – temperature and rainfall – for trends, using state-space models that allowed us to account for seasonal effects.

This code chunk reads in the data and prepares the temperature variable as a time series.

```
my_data <- read.csv("WagtailWeather.csv")
names(my_data)

## [1] "Year"    "Season"   "Temp"     "Rain"

temp_1 <- my_data$Temp
temp_2 <- as.matrix(temp_1)
temp = ts(temp_2, start = 1976, frequency=4)
```

Then, we fit a state-space model to these data.

State Space Model for temperature

```
library(dlm)
library(Hmisc)

fit.st <- StructTS (temp, type = "BSM")

buildFun <- function(x){
  dlmTemp <- dlmModPoly(2) + dlmModSeas(4)
  diag(W(dlmTemp))[2:3] <- exp(x[1:2])
  V(dlmTemp) <- exp(x[3])
  return(dlmTemp)
}

buildFun2 <- function(x){
  dlmTemp <- dlmModPoly(2) + dlmModSeas(4)
  diag(W(dlmTemp))[2] <- exp(x[1])
  V(dlmTemp) <- exp(x[2])
  return(dlmTemp)
}

fit <- dlmMLE(temp, parm = rep(0, 3), build = buildFun)
fit2 <- dlmMLE(temp, parm = rep(0, 2), build = buildFun2)
```

```

dlmTemp <- buildFun(fit$par)
drop(V(dlmTemp))

## [1] 0.1541838
diag(W(dlmTemp)) [2:3]

## [1] 4.543804e-04 2.542649e-10
tempSmooth <- dlmSmooth(temp, mod = dlmTemp)
sm <- ts(tempSmooth$s[-1], start = 1977, frequency = 4)

```

Plot the model fit:

```

par(mfrow=c(2,1), mar=c(0,4,1,1), oma=c(3,0,0,0))
plot(temp, type = "o", cex.lab = 1.5, cex.axis = 1.5, las = 1, xlim = c(1975.5, 2000),
     ylab = "Temperature", axes=F)
minor.tick(nx = 5, ny = 0, tick.ratio = 0.5)
axis(2, las=1)

lines(tempSmooth$s[, 1], col = 2, type = "l", pch = 16, cex = 0.7)
abline(h = mean(temp), col = "grey")

# ----- confidence intervals -----
U.S <- tempSmooth$U.S
D.S <- tempSmooth$D.S

v <- dlmSvd2var(U.S, D.S)
vars <- c()

for (i in 1:(length(temp) + 1))
{ vars[i] <- v[[i]][1, 1]
}

pl <- tempSmooth$s[, 1] + qnorm(0.025, sd = sqrt(vars))
pu <- tempSmooth$s[, 1] + qnorm(0.975, sd = sqrt(vars))

lines(pl, lty = 2)
lines(pu, lty = 2)

plot(tempSmooth$s[, 2], ylab="Slope", xlab="Year", axes=F, ylim = c(-0.15, 0.2),
      xlim = c(1975.5, 2000), type = "l")
abline(h=0, col="grey")
minor.tick(nx = 5, ny = 0, tick.ratio = 0.5)
axis(1, at=c(1976:2000))
axis(2, las=1)

for (i in 1:(length(temp) + 1))
{ vars[i] <- v[[i]][2, 2]
}

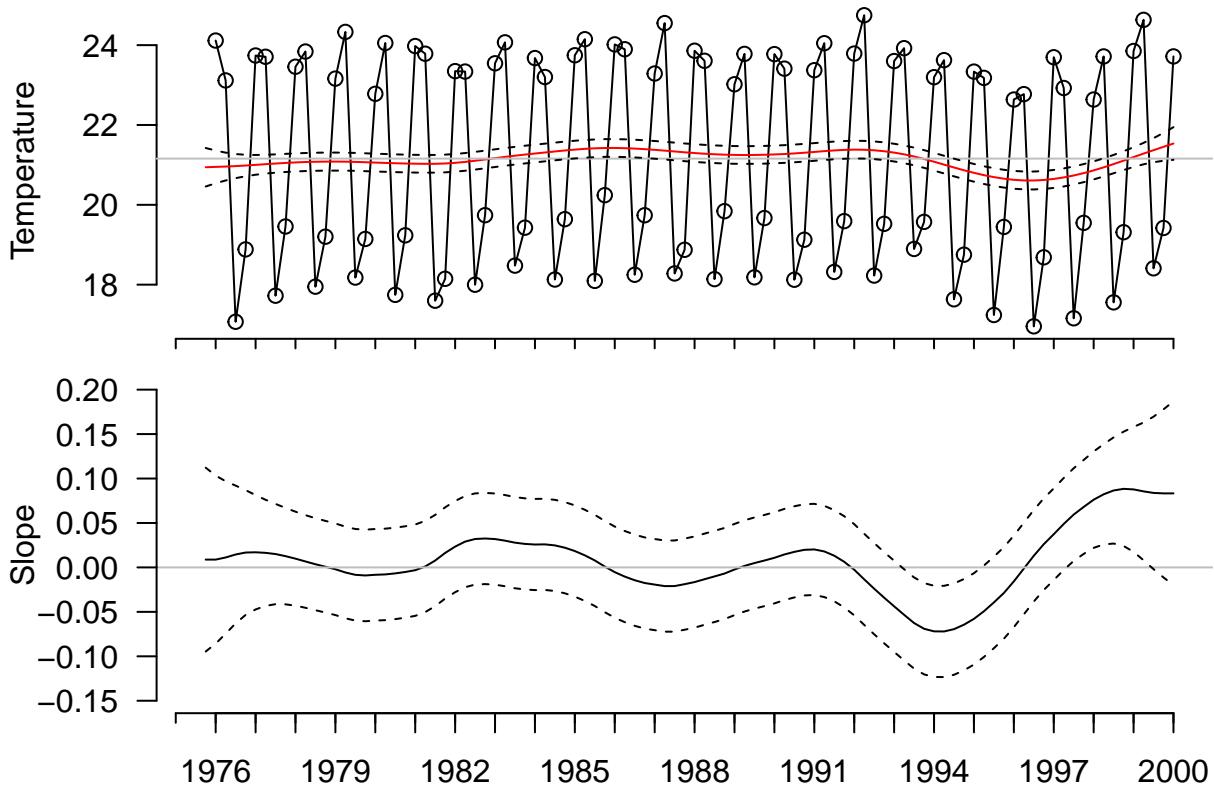
pl <- tempSmooth$s[, 2] + qnorm(0.025, sd = sqrt(vars))
pu <- tempSmooth$s[, 2] + qnorm(0.975, sd = sqrt(vars))

```

```

lines(pl, lty = 2)
lines(pu, lty = 2)

```



State Space Model for rainfall

```

rain_1 <- my_data$Rain
rain_2 <- as.matrix(rain_1)
rain <- ts(rain_2, start=1976, frequency=4)

fit.st <- StructTS (rain, type = "BSM")

buildFun <- function(x){
  dlmTemp <- dlmModPoly(2) + dlmModSeas(4)
  diag(W(dlmTemp)) [2:3] <- exp(x[1:2])
  V(dlmTemp) <- exp(x[3])
  return(dlmTemp)
}

buildFun2 <- function(x){
  dlmTemp <- dlmModPoly(2) + dlmModSeas(4)
  diag(W(dlmTemp)) [2] <- exp(x[1])
  V(dlmTemp) <- exp(x[2])
  return(dlmTemp)
}

```

```

fit <- dlmMLE(rain, parm = rep(0, 3), build = buildFun)
fit2 <- dlmMLE(rain, parm = rep(0, 2), build = buildFun2)

fit <- fit2

dlmTemp <- buildFun2(fit$par)
drop(V(dlmTemp))

## [1] 12149.89
diag(W(dlmTemp)) [2:3]

## [1] 2.04632 1.00000
tempSmooth <- dlmSmooth(rain, mod = dlmTemp)
sm <- ts(tempSmooth$s[-1], start = 1977, frequency = 4)

```

Plotting the model fit:

```

par(mfrow=c(2, 1), mar=c(0,4,1,1), oma=c(3,0,0,0))

plot(rain, type = "o", cex.lab = 1.5, cex.axis = 1.5, las = 1,
      ylab ="Rain [mm]", axes=F, xlim = c(1975.5, 2000))
minor.tick(nx = 5, ny = 0, tick.ratio = 0.5)
axis(2, las=1)

lines(tempSmooth$s[, 1], col = 2, type = "l", pch = 16, cex = 0.7)
abline(h = mean(rain), col = "grey")

# ----- confidence intervals -----
U.S <- tempSmooth$U.S
D.S <- tempSmooth$D.S

v <- dlmSvd2var(U.S, D.S)
vars <- c()

for (i in 1:(length(rain) + 1))
{ vars[i] <- v[[i]][1, 1]
}

pl <- tempSmooth$s[, 1] + qnorm(0.025, sd = sqrt(vars))
pu <- tempSmooth$s[, 1] + qnorm(0.975, sd = sqrt(vars))

lines(pl, lty = 2)
lines(pu, lty = 2)

## slope
plot(tempSmooth$s[, 2], ylab="Slope", xlab="Year", axes=F, xlim = c(1975.5, 2000),
      ylim = c(-20, 20))
abline(h=0, col="grey")
minor.tick(nx = 5, ny = 0, tick.ratio = 0.5)
axis(2, las=1)

for (i in 1:(length(rain) + 1))
{ vars[i] <- v[[i]][2, 2]
}

```

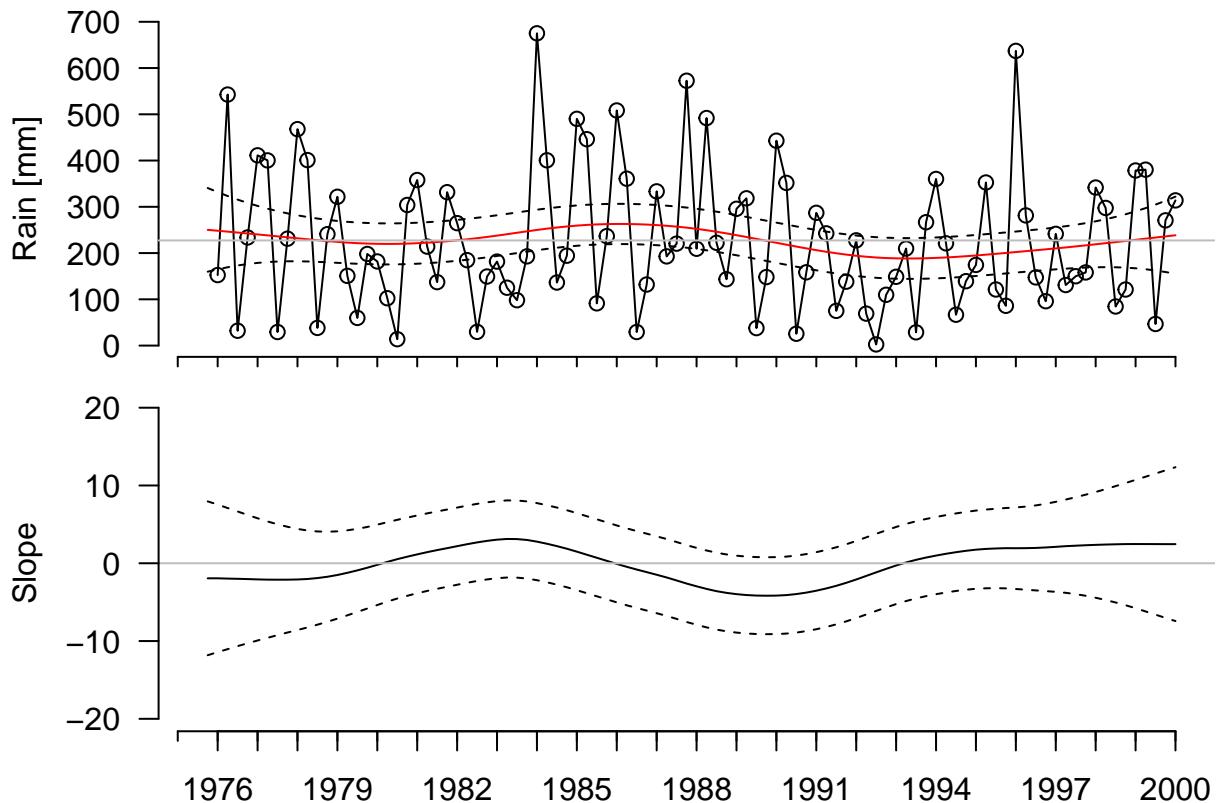
```

}

pl <- tempSmooth$s[, 2] + qnorm(0.025, sd = sqrt(vars))
pu <- tempSmooth$s[, 2] + qnorm(0.975, sd = sqrt(vars))

lines(pl, lty = 2)
lines(pu, lty = 2)
axis(1, at=c(1976:2000))

```



Mass trends

Then, we look at trends in the body mass of mountain wagtails. We first see whether there is any evidence for a linear change in mass over years and whether mass changes seasonally.

```

temp <- read.csv("WagtailMass.csv")

temp$time <- temp$year + (temp$season-1)/4 # time is now continuous, in quarters

temp$season <- as.factor(temp$season)

out.lm1 <- lm(mass ~ I(time-mean(time)) * season, data = temp)
summary(out.lm1) # there is a clear year effect

##
## Call:

```

```

## lm(formula = mass ~ I(time - mean(time)) * season, data = temp)
##
## Residuals:
##   Min     1Q Median     3Q    Max 
## -5.2364 -0.7344 -0.0462  0.8109  3.2804 
## 
## Coefficients:
##                               Estimate Std. Error t value Pr(>|t|)    
## (Intercept)               19.739398  0.101340 194.784 < 2e-16 ***
## I(time - mean(time))      -0.044139  0.014923 -2.958  0.00323 **  
## season2                   -0.190895  0.131751 -1.449  0.14792  
## season3                   0.095626  0.152923  0.625  0.53201  
## season4                   0.013339  0.169225  0.079  0.93720  
## I(time - mean(time)):season2 0.027479  0.018454  1.489  0.13703  
## I(time - mean(time)):season3 -0.009618  0.022241 -0.432  0.66557  
## I(time - mean(time)):season4 -0.010075  0.026492 -0.380  0.70387  
## ---                        
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1 
## 
## Residual standard error: 1.219 on 566 degrees of freedom
## Multiple R-squared:  0.05705,   Adjusted R-squared:  0.04539 
## F-statistic: 4.892 on 7 and 566 DF,  p-value: 2.236e-05
anova(out.lm1)      # but no evidence for mass to depend on season

## Analysis of Variance Table
##
## Response: mass
##                               Df Sum Sq Mean Sq F value Pr(>F)    
## I(time - mean(time))       1  35.17  35.169 23.6735 1.482e-06 ***
## season                      3    7.84   2.614  1.7595   0.1538  
## I(time - mean(time)):season 3    7.86   2.620  1.7639   0.1529  
## Residuals                  566 840.84   1.486  
## ---                        
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1 
out.lm2 <- lm(mass ~ I(time-mean(time)) + season, data = temp)
summary(out.lm2) # removing season-time interaction does not change the time effect

## 
## Call:
## lm(formula = mass ~ I(time - mean(time)) + season, data = temp)
## 
## Residuals:
##   Min     1Q Median     3Q    Max 
## -5.1087 -0.7040 -0.0137  0.8387  3.1796 
## 
## Coefficients:
##                               Estimate Std. Error t value Pr(>|t|)    
## (Intercept)               19.749029  0.100668 196.180 < 2e-16 ***
## I(time - mean(time))     -0.034705  0.007319 -4.741 2.68e-06 ***  
## season2                  -0.206789  0.131169 -1.577  0.115    
## season3                  0.096588  0.152229  0.634   0.526    
## season4                  -0.051253  0.159658 -0.321   0.748    
## ---                        

```

```

## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 1.221 on 569 degrees of freedom
## Multiple R-squared:  0.04823,   Adjusted R-squared:  0.04154
## F-statistic: 7.209 on 4 and 569 DF,  p-value: 1.148e-05
out.lm3 <- lm(mass ~ time, data = temp)
summary(out.lm3) # removing season does not change the time effect

##
## Call:
## lm(formula = mass ~ time, data = temp)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -5.2486 -0.6900 -0.0245  0.8340  3.1944
##
## Coefficients:
##             Estimate Std. Error t value Pr(>|t|)
## (Intercept) 89.163263  14.337011   6.219 9.66e-10 ***
## time        -0.034905   0.007203  -4.846 1.62e-06 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 1.224 on 572 degrees of freedom
## Multiple R-squared:  0.03944,   Adjusted R-squared:  0.03776
## F-statistic: 23.49 on 1 and 572 DF,  p-value: 1.623e-06
coef(out.lm3)[2] * 10/mean(temp$mass)  # percent decline in mass per decade

##          time
## -0.01773335

1+coef(out.lm3)[2] * 10/mean(temp$mass)  # percent decline in mass per decade

##
##          time
## 0.9822667

(1+coef(out.lm3)[2] * 10/mean(temp$mass))^0.75  # change in metabolic rate

##          time
## 0.9866703

```

Then we want to separate the among-individual and within-individual effects, following van de Pol and Wright 2009, Animal Behaviour.

```

# calculate mean time per individual
meantime <- aggregate(temp$time, by = list(temp$ringno), FUN=mean)
colnames(meantime) <- c("ringno", "meantime")

# add mean time to data frame
temp <- merge(temp, meantime, by.x = "ringno", by.y = "ringno", all.x=T)
temp$dtime <- temp$time - temp$meantime  # difference in time within individual

library(nlme)

# almost the entire effect is due to heavier individuals being replaced by lighter individuals
out.lme1 <- lme(mass ~ dtime + meantime, random =~1|ringno, data=temp)

```

```

summary(out.lme1)

## Linear mixed-effects model fit by REML
##  Data: temp
##        AIC      BIC    logLik
##   1748.062 1769.799 -869.0311
##
## Random effects:
##   Formula: ~1 | ringno
##             (Intercept) Residual
##   StdDev:     1.038572 0.7699628
##
## Fixed effects: mass ~ dtime + meantime
##                  Value Std.Error DF t-value p-value
## (Intercept) 73.03214 20.772130 287 3.515871 0.0005
## dtime       -0.00857  0.026532 287 -0.323024 0.7469
## meantime    -0.02686  0.010434 284 -2.574271 0.0106
##
## Correlation:
##          (Intr) dtime
## dtime      0
## meantime -1      0
##
## Standardized Within-Group Residuals:
##      Min      Q1      Med      Q3      Max
## -3.79528290 -0.46563894 -0.04904805  0.46075842  3.71350367
##
## Number of Observations: 574
## Number of Groups: 286
# proportion of trend explained by among-individual effect
fixef(out.lme1)[3]/sum(fixef(out.lme1)[2:3])

```

```

## meantime
## 0.7581056

```

Then look at detailed trend using GAM

```

library(mgcv)

## This is mgcv 1.8-23. For overview type 'help("mgcv-package")'.
gam.mass <- gam(mass ~ s(time), data = temp)

```

Finally plot it all:

```

plot(mass~time, data=temp, xlab="Year", ylab="Mass [g]", axes=F, col="grey")
axis(1)
axis(2,las=1)
abline(out.lm3, lwd=2)
x <- seq(min(temp$meantime), max(temp$meantime), length=50)
lines(x,fixef(out.lme1)[1] + fixef(out.lme1)[3]*x,lty=2, col="blue", lwd=2)
od<-order(temp$time)
lines(temp$time[od], predict(gam.mass)[od], lty=3, col="red", lwd=3)

```

