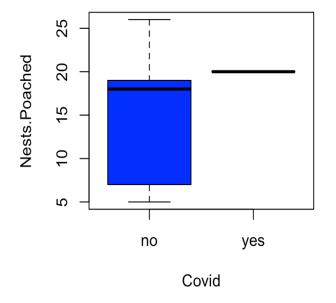
```
> m1<-glmmPQL(Visits~Covid,random=~1|Type,data=CV,family=quasipoisson)</pre>
iteration 1
iteration 2
> summary(m1)
Linear mixed-effects model fit by maximum likelihood
 Data: CV
 AIC BIC logLik
                                                                   800
  NA NA
              NA
Random effects:
                                                                   900
 Formula: ~1 | Type
        (Intercept) Residual
StdDev: 0.04106479 8.116669
                                                                   400
Variance function:
                                                                   200
 Structure: fixed weights
 Formula: ~invwt
Fixed effects: Visits ~ Covid
                Value Std.Error DF t-value p-value
(Intercept) 6.276294 0.1692661 5 37.07945 0.0000
                                                                      2017.0
                                                                                2018.0
                                                                                           2019.0
                                                                                                     2020.0
Covidyes
            -1.179517 0.5441911 5 -2.16747 0.0824
Correlation:
         (Intr)
                                                                                       Year
Covidves -0.299
Standardized Within-Group Residuals:
                                                                          69% drop in park visits
                   01
       Min
                             Med
                                         Q3
                                                   Max
-1.1993669 -0.7775371 -0.2429437 0.9684828 1.4189836
Number of Observations: 8
Number of Groups: 2
```

> round(((exp(6.276294-1.179517)-exp(6.276294))/exp(6.276294))*100)

[1] -69

```
> m1<-gls(Nests.Poached~Covid,data=NP)</pre>
> summary(m1)
Generalized least squares fit by REML
  Model: Nests.Poached ~ Covid
  Data: NP
       AIC
                BIC
                       logLik
  36.36206 34.52094 -15.18103
Coefficients:
            Value Std.Error t-value p-value
               15 3.937004 3.810004 0.0189
(Intercept)
Covidyes
                5 9.643651 0.518476 0.6315
 Correlation:
         (Intr)
Covidyes -0.408
Standardized residuals:
       Min
                   Q1
                             Med
                                         Q3
                                                   Max
-1.1359237 -0.6815542 0.1703886 0.4259714 1.2495160
Residual standard error: 8.803408
Degrees of freedom: 6 total; 4 residual
> round((5/15)*100)
[1] 33
```



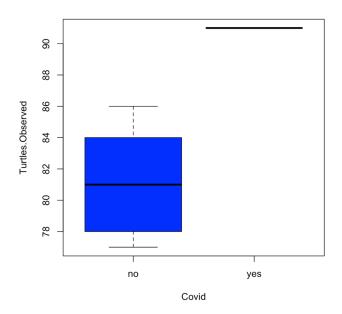
33% increase in poaching

```
> m1<-gls(Turtles.Observed~Covid,data=T0)</pre>
> summary(m1)
Generalized least squares fit by REML
  Model: Turtles.Observed ~ Covid
  Data: TO
       AIC
                BIC
                       logLik
  29.71234 27.87122 -11.85617
Coefficients:
            Value Std.Error t-value p-value
(Intercept) 81.2 1.714643 47.35680
                                        0.00
Covidyes
              9.8 4.200000 2.33333
                                        0.08
 Correlation:
         (Intr)
Covidyes -0.408
Standardized residuals:
                     Q1
                                Med
        Min
                                             Q3
                                                        Max
-1.09544512 -0.63900965 -0.02608203 0.54772256 1.25193727
Residual standard error: 3.834058
```

Degrees of freedom: 6 total; 4 residual

> round((9.8/81.2)*100)

[1] 12



12% increase in turtles observed, but with > effort