# Simulating and estimating the hierarchical lengths example from Bence

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# R code for simulating a mixed effects model: lengths.R

#### lengths.R cont'd

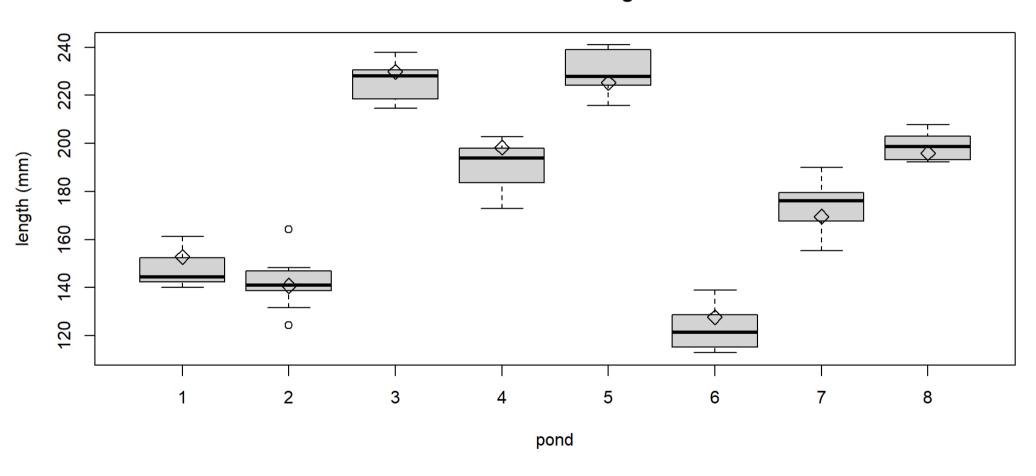
```
1  # step 1, calculate pond mean lengths from hyperprior:
2  set.seed(31)
3  mu pond = rnorm(n pond, mu, sd among) # average length in each lake
```

#### lengths.R cont'd

```
1 # step 2, calculate fish lengths | on mu_pond:
2 y_obs = rep(NA, n_pond * n_fish)
3
4 # loop through data, generate a random fish length | mu_pond
5 for (i in 1:length(y_obs)) {
6  y_obs[i] = rnorm(1, mu_pond[pond[i]], sd_within)
7 }
```

#### Look at the simulated data

#### distribution of lengths



# TMB code for estimating the lengths model: lengths.cpp

```
1 #include <TMB.hpp>
 2 template<class Type>
   Type objective function<Type>::operator() ()
 4 {
    DATA VECTOR(y obs); // observed data
    DATA INTEGER (n pond);
     DATA IVECTOR (pond);
                             // integer vector indicating pond
 8
                        // average length among lakes
9
     PARAMETER (ln mu);
     PARAMETER(ln sd among); // sd among ponds
10
     PARAMETER(ln sd within); // sd within ponds
11
     PARAMETER VECTOR(eps); // random effects - deviations from mu
12
13
     int n obs = y obs.size();
14
15
     Type mu = exp(ln mu);
     Type sd among = exp(ln sd among);
16
     Type sd within = exp(ln sd within);
17
18
19
     Type jnll = 0.0;
20
21
     // Pr(random coefficients)
     for (int i = 0; i < n pond; i++) {</pre>
       jnll -= dnorm(eps(i), Type(0.0), sd among, true);
23
24
25
     // more code below
```

## TMB code: lengths.cpp

```
// continued from previous slide
// Pr(data conditional on fixed and random effect values)
for(int i = 0; i < n_obs; i++) {
    jnll -= dnorm(y_obs(i), mu + eps(pond(i)), sd_within, true);
}
return jnll;
}</pre>
```

### R code: lengths.R

```
1 library (TMB)
 2 compile("lengths.cpp")
[1] 0
 1 dyn.load(dynlib("lengths"))
 3 	 data = list(
 4 	 y 	 obs = y 	 obs,
 5 n \text{ pond} = n \text{ pond}
    pond = pond - 1 + look here, this is important
 7
 8
   parameters = list(
10
    ln mu = log(150),
11 ln sd among = log(50),
12 \ln sd within = \log(50),
eps = rep(0, n pond)
14 )
15
16 obj = MakeADFun(data, parameters, random = "eps", DLL = "lengths")
```

### Testing objective function

```
1 obj$fn() # return the objective f(x) value
Optimizing tape... Done
iter: 1 value: 429.3855 mgc: 0.3160354 ustep: 1
iter: 2 mgc: 5.182486e-17

[1] 400.3294
attr(,"logarithm")
[1] TRUE
```

## Testing gradients

```
1 obj$gr() # examine par gradients
iter: 1 mgc: 5.182486e-17
Matching hessian patterns... Done
outer mgc: 69.99745
[1] -12.631046    1.480762    69.997450
```

#### Run the optimization:

```
1 opt = nlminb(obj$par, obj$fn, obj$gr)
iter: 1 mgc: 5.182486e-17
iter: 1 mgc: 5.182486e-17
outer mgc: 69.99745
iter: 1 value: 356.1635 mgc: 0.8939608 ustep: 1
iter: 2 mgc: 6.782769e-16
iter: 1 value: 2960.189 mgc: 1329.285 ustep: 1
iter: 2 mgc: 5.255102e-13
iter: 1 mgc: 6.782769e-16
outer mgc: 56.6637
iter: 1 value: 605.3189 mgc: 2.211058 ustep: 1
iter: 2 mgc: 2.735832e-14
iter: 1 value: 338.4739 mgc: 0.02171076 ustep: 1
iter: 2 mgc: 1.028691e-15
iter: 1 mgc: 1.028691e-15
outer mgc: 44.02805
```

#### Get standard deviations

```
1 sdr = sdreport(obi)
iter: 1 mgc: 2.669739e-15
outer mgc: 2.076325e-05
iter: 1 value: 322.2081 mgc: 0.02397546 ustep: 1
iter: 2 mgc: 2.359224e-15
outer mgc: 0.1897302
iter: 1 value: 322.2081 mgc: 0.0239515 ustep: 1
iter: 2 mgc: 1.852685e-15
outer mgc: 0.1891672
iter: 1 value: 322.208 mgc: 8.261608e-05 ustep: 1
iter: 2 mgc: 1.132774e-15
outer mgc: 0.0158112
iter: 1 value: 322.2079 mgc: 8.278147e-05 ustep: 1
iter: 2 mgc: 1.619798e-15
outer mgc: 0.01583635
iter: 1 value: 322.216 mgc: 8.261608e-05 ustep: 1
```

#### Get standard deviations

## **Check diagnostics**

```
1 # check gradients and make sure pdHess
2 opt$convergence # 0 is good

[1] 0
1 final_gradient <- obj$gr(opt$par)

iter: 1 mgc: 2.669739e-15
outer mgc: 2.076325e-05

1 if (any(abs(final_gradient) > 0.001) || sdr$pdHess == FALSE) {
    message("Model did not converge: check results")
    } else {
        message("Model diagnostics consistent with convergence")
    }
}
```

#### Plot fits vs. data

#### distribution of lengths

