Homework - Week 3

Vasco Brazão

```
library(rethinking)
library(tidyverse)

set.seed(22012022)

data(foxes)

d <- foxes</pre>
```

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```
d1 <- d %>%
  dplyr::mutate(
    A = rethinking::standardize(area),
    F = rethinking::standardize(avgfood)
)
```

Since there is no back-door at all, we can simply regress Food on Area to get the total (and, in this case, also direct) causal effect of A on F.

```
\begin{aligned} \mathbf{F}_i &\sim \mathrm{Normal}(\mu_i, \sigma) \\ \mu_i &= \alpha + \beta \mathbf{A}_i \\ \alpha &\sim \mathrm{Normal}(0, 0.2) \\ \beta &\sim \mathrm{Normal}(0, 1) \\ \sigma &\sim \mathrm{Exponential}(1) \end{aligned}
```

```
m1 <- rethinking::quap(
    alist(
        F ~ dnorm(mu, sigma),
        mu <- a + b*A,
        a ~ dnorm(0, 0.2),
        b ~ dnorm(0, 1),
        sigma ~ dexp(1)
    ),
    data = d1
)
precis(m1)</pre>
```

```
## mean sd 5.5% 94.5%

## a 7.843048e-08 0.04230780 -0.06761596 0.06761611

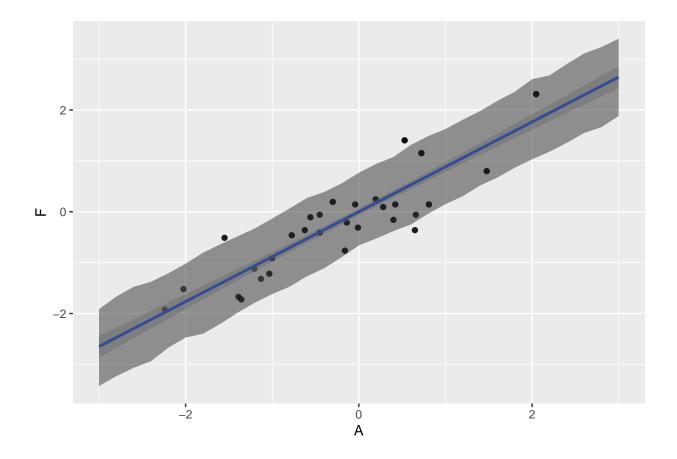
## b 8.814382e-01 0.04343473 0.81202108 0.95085526

## sigma 4.662197e-01 0.03051689 0.41744782 0.51499160
```

From the marginal distribution of β , we conclude that each standard deviation increase in Area increases Food by .88 SD.

For a plot of the whole thing:

```
# mu and 89% PI for mu across all areas in the data
area.seq <- tibble::tibble(</pre>
  A = seq(-3, 3, by = 0.2)
mu1 <- rethinking::link(m1, data = area.seq)</pre>
mu1.mean <- apply(mu1, 2, mean)</pre>
mu1.pi <- apply(mu1, 2, rethinking::PI, prob = 0.89)</pre>
mu1.tibble <- area.seq %>%
  dplyr::mutate(
    mu1.mean = mu1.mean,
    mu1.lower = mu1.pi[1,],
    mu1.upper = mu1.pi[2,]
  )
# simulating weights to calculate 89% prediction interval
sim.weight <- rethinking::sim(m1, data = list(A = area.seq$A))
weight.PI <- apply(sim.weight, 2, rethinking::PI, prob = .89)</pre>
pred.tibble <- area.seq %>%
  dplyr::mutate(
    weight.lower = weight.PI[1,],
    weight.upper = weight.PI[2,],
  )
d1 %>%
  ggplot2::ggplot(ggplot2::aes(x = A, y = F)) +
  ggplot2::geom_point(alpha = 0.5) +
  ggplot2::geom_smooth(aes(x = A, y = mu1.mean, ymin = mu1.lower, ymax = mu1.upper), data = mu1.tibble,
  ggplot2::geom\_ribbon(aes(x = A, y = NULL, ymin = weight.lower, ymax = weight.upper), data = pred.tibb
```



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```
d2 <- d %>%
  dplyr::mutate(
    A = rethinking::standardize(area),
    F = rethinking::standardize(avgfood),
    W = rethinking::standardize(weight),
    G = rethinking::standardize(groupsize)
)
```

For the total causal effect, we can simply regress weight on food.

$$W_i \sim \text{Normal}(\mu_i, \sigma)$$
$$\mu_i = \alpha + \beta F_i$$
$$\alpha \sim \text{Normal}(0, 0.2)$$
$$\beta \sim \text{Normal}(0, 1)$$
$$\sigma \sim \text{Exponential}(1)$$

```
m2 <- rethinking::quap(
    alist(
        W ~ dnorm(mu, sigma),
        mu <- a + b*F,
        a ~ dnorm(0, 0.2),
        b ~ dnorm(0, 1),
        sigma ~ dexp(1)
    ),
    data = d2
)

precis(m2)</pre>
```

```
## mean sd 5.5% 94.5%

## a -1.027080e-06 0.08360010 -0.1336101 0.1336081

## b -2.482821e-02 0.09203231 -0.1719136 0.1222572

## sigma 9.911429e-01 0.06465840 0.8878063 1.0944795
```

Somehow adding food seems to have a rather small effect on the weight of the foxes.

I would speculate that more food leads to larger groups so that each individual fox ends up eating about the same anyway.

Let's check the direct effect of food on weight by adding groupsize as a predictor.

```
W_i \sim \text{Normal}(\mu_i, \sigma)
\mu_i = \alpha + \beta_F F_i + \beta_G G_i
\alpha \sim \text{Normal}(0, 0.2)
\beta_j \sim \text{Normal}(0, 1)
\sigma \sim \text{Exponential}(1)
```

```
m2.2 <- rethinking::quap(
    alist(
        W ~ dnorm(mu, sigma),
        mu <- a + bF*F + bG*G,
        a ~ dnorm(0, 0.2),
        bF ~ dnorm(0, 1),
        bG ~ dnorm(0, 1),
        sigma ~ dexp(1)
    ),
    data = d2
)

precis(m2.2)</pre>
```

```
## mean sd 5.5% 94.5%

## a -9.951017e-06 0.07995525 -0.1277939 0.1277740

## bF 5.916044e-01 0.19540629 0.2793074 0.9039014

## bG -6.890407e-01 0.19540789 -1.0013403 -0.3767412

## sigma 9.394854e-01 0.06133642 0.8414580 1.0375129
```

Indeed, for each level of groupsize, more food seems to imply more weight. And symmetrically, once we know the amount of food, higher group sizes are indicative of lower individual weight.

Our DAG suggests that food comes "first", meaning that the story we associate with both models is that higher food leads directly to higher weight, but it also leads to larger groups. Since more individuals then all share the same food, that has a negative effect on the weight, balancing out. Perhaps groups of foxes grow in a sort of equilibrium, where they become more numerous if food allows but only as long as each individual maintains a minimum level of weight.

But that's for urban ecologists to theorize.