Bayesian Workflow

Maximizing the odds of building a useful model!

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Bayesian Workflow













Model height as a function of weight:

```
data {
    int num_people;
    vector<lower=0>[num_people] weights;
    vector<lower=0> heights[num_people];
}
```



Construct mathematical model you think generated the data:

height ~
$$\mathcal{N}(\beta * \text{weight} + \alpha, \sigma)$$

In Stan, we'd now write the parameters of this model:

```
parameters {
   real beta;
   real alpha;
   real<lower=0> sigma;
}
```



Construct mathematical model you think generated the data:

height ~
$$\mathcal{N}(\beta * \text{weight} + \alpha, \sigma)$$

And then write the likelihood:

```
model {
    heights ~ normal(beta * weights + alpha, sigma);
}
```

height ~ $\mathcal{N}(\beta * \text{weight} + \alpha, \sigma)$

Think about reasonable priors for your parameters:

- Beta measures the association between weight and height, in cm/kg
- Alpha is the intercept, or average height for someone with no weight (not a
 particularly useful number on its own)
- Sigma is the standard deviation capturing un-modeled variation in the population

```
In Stan:
```

```
model {
   heights ~ normal(beta * weights + alpha, sigma);
   beta ~ normal(0, 10); // cm/kg
   alpha ~ normal(50, 50); // avg cm for 0 kg
   sigma ~ normal(0, 5); // variation from average
}
```

Sanity check:

- 1. Draw parameter values from priors
- 2. Generate data based on those parameter values
- 3. Fit model to generated data
- 4. Check fit is reasonable

```
generated quantities {
  real<lower=0> heights[N];
  real beta = normal_rng(0, 10);
  real alpha = normal_rng(50, 50);
  real sigma = fabs(normal_rng(0, 5));
  for (n in 1:N)
     heights[n] = normal_rng(beta * weights[n] + alpha, sigma);
}
```



Warning messages:

1: There were 13 divergent transitions after warmup. Increasing adapt_delta above 0.8 may help. See <u>http://mc-</u> <u>stan.org/misc/warnings.html#divergent-transitions-after-warmup</u> 2: There were 70 transitions after warmup that exceeded the maximum treedepth. Increase max_treedepth above 10. See <u>http://mcstan.org/misc/warnings.html#maximum-treedepth-exceeded</u> 3: There were 4 chains where the estimated Bayesian Fraction of Missing Information was low. See <u>http://mcstan.org/misc/warnings.html#bfmi-low</u>

4: Examine the pairs() plot to diagnose sampling problems





For each sample from parameter space, generate some fake measurements, and see how they match up against the real measurements.

```
generated quantities {
   real h_ppc[N];
   for (n in 1:N)
      h_ppc[n] = normal_rng(beta * weights[n] + alpha, sigma);
}
```







Use these tools to compare predictive distributions and other quantities of interest among models.

Iterate!

