

The Normal Model

Professor Rodrigo Targino

Announcements

- Reading: Section 5.3.3 and 5.3.4

The Normal Distribution

- One of the most utilized probability models in data analysis
- Central Limit Theorem
- Separate parameters for the mean and the variance (intuitive)

Normal Distribution

- Symmetric with mode = median = mean = μ
- Approximately 95% of the population lies within two standard deviations of the mean
- Density:

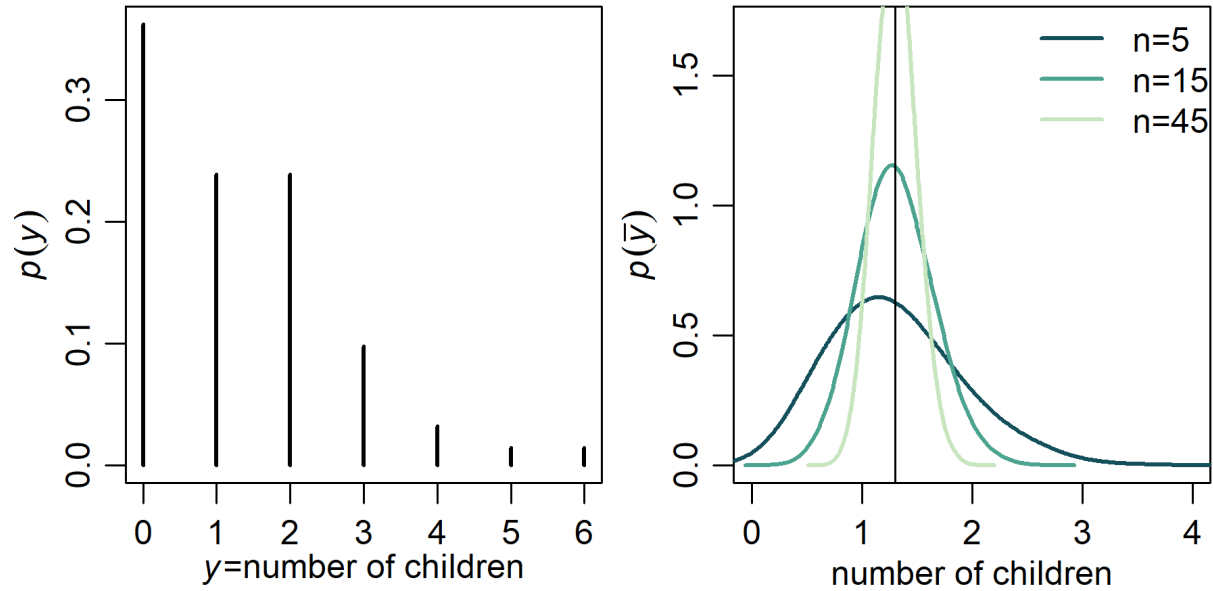
$$p(y|\mu, \sigma^2) = \frac{1}{\sqrt{2\pi\sigma^2}} e^{-\frac{1}{2}\left(\frac{y-\mu}{\sigma}\right)^2}, \quad -\infty < y < \infty$$

- $X \sim N(\mu_x, \sigma_x^2)$ and $Y \sim N(\mu_y, \sigma_y^2)$ with X and Y independent then

$$aX + bY \sim N(a\mu_x + b\mu_y, a^2\sigma_x^2 + b^2\sigma_y^2)$$

- In R: `dnorm`, `rnorm`, `pnorm`, `qnorm`.
 - Warning: the argument to the `norm` functions in R is σ not σ^2 !

The Central Limit Theorem



$$\text{CLT: } \bar{y} \approx N(E[Y], \text{Var}[Y]/n)$$

Bayesian inference in the normal model

- Assume $y_1, \dots, y_n \sim N(\mu, \sigma^2)$ with σ^2 a known constant
- Lets start with a non-informative, improper prior: $p(\mu) \propto \text{const}$
- What is the posterior distribution $p(\mu \mid y_1, \dots, y_n, \sigma^2)$?

Bayesian inference in the normal model

Bayesian inference in the normal model

Bayesian inference in the normal model

- Assume $y_1, \dots, y_n \sim N(\mu, \sigma^2)$ with σ^2 a known constant
- The normal prior distribution is conjugate for μ in the normal sampling model
- Sampling distribution, prior distribution and posterior distribution are all normal.
- Assume the prior is $p(\mu) \sim N(\mu_0, \tau^2)$
- What are the parameters of the posterior $p(\mu \mid y_1, \dots, y_n, \sigma^2)$?

Bayesian inference in the normal model

Bayesian inference in the normal model

A conjugate prior for the normal likelihood

- The normal distribution is conjugate for the normal likelihood
 - Often called the "normal-normal model"
- $Y_i \sim N(\mu, \sigma^2)$ and $\mu \sim N(\mu_0, \tau^2)$ implies that the posterior distribution $p(\mu \mid y)$ is also normally distributed:

$$\mu \mid Y \sim N(\mu_n, \tau_n^2)$$

$$\text{where } \mu_n = \frac{\frac{1}{\tau^2}\mu_0 + \frac{n}{\sigma^2}\bar{y}}{\frac{1}{\tau^2} + \frac{n}{\sigma^2}} \text{ and } \tau_n^2 = \frac{1}{\frac{1}{\tau^2} + \frac{n}{\sigma^2}}$$

The posterior mean and pseudo-counts

$$\begin{aligned}\mu_n &= \frac{\frac{1}{\tau^2}}{\frac{1}{\tau^2} + \frac{n}{\sigma^2}} \mu_0 + \frac{\frac{n}{\sigma^2}}{\frac{1}{\tau^2} + \frac{n}{\sigma^2}} \bar{y} \\ &= (1 - w) \mu_0 + w \bar{y}\end{aligned}$$

$$\text{where } w = \frac{\frac{n}{\sigma^2}}{\frac{1}{\tau^2} + \frac{n}{\sigma^2}}$$

Can we think about the normal prior parameters in terms of pseudo-counts?

The posterior mean and pseudo-counts

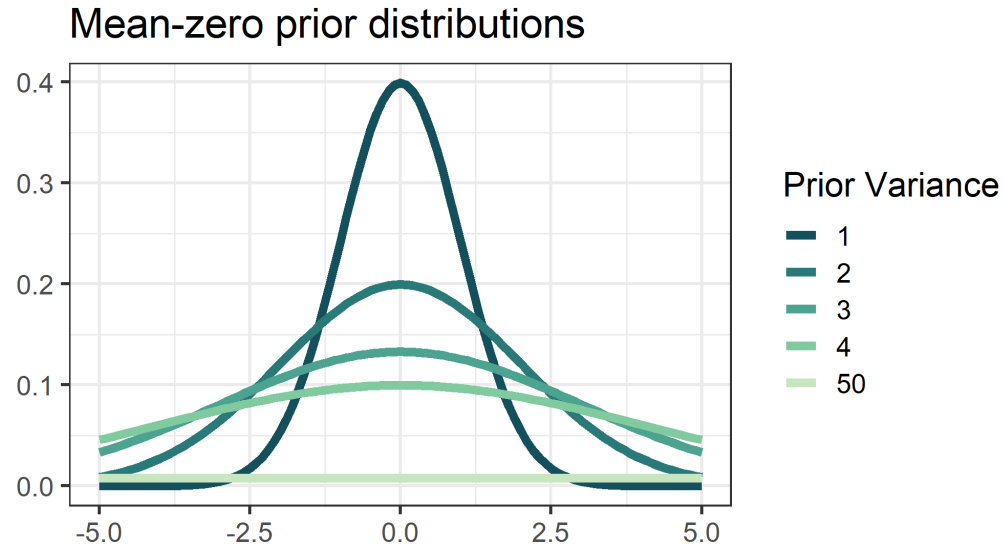
$$\begin{aligned}\mu_n &= \frac{\frac{1}{\tau^2}}{\frac{1}{\tau^2} + \frac{n}{\sigma^2}} \mu_0 + \frac{\frac{n}{\sigma^2}}{\frac{1}{\tau^2} + \frac{n}{\sigma^2}} \bar{y} \\ &= (1 - w) \mu_0 + w \bar{y}\end{aligned}$$

where $w = \frac{\frac{n}{\sigma^2}}{\frac{1}{\tau^2} + \frac{n}{\sigma^2}}$

- Let's reparameterize: $\tau^2 = \frac{\sigma^2}{\kappa_0}$
- Then: the posterior variance is $\frac{\sigma^2}{\kappa_0 + n}$
- And: $(1 - w) = \frac{\kappa_0}{\kappa_0 + n}$
- κ_0 are the prior counts and μ_0 is the prior sample average.

Conjugate prior with increasing variance

```
## Warning: Using `size` aesthetic for lines was deprecated in ggplot2 3.4.  
## i Please use `linewidth` instead.
```



Bayes Estimators

Estimators: Bayes / Frequentist Unification

- Bayesian inference provides a straightforward procedure for producing estimators given your prior beliefs.
 1. Compute posterior distribution
 2. Summarize the posterior distribution with a point estimator (e.g. posterior mean or posterior mode) and a probability interval
- Frequentists provide tools for evaluating the sampling properties of an estimator.
 - Bias, variance and MSE of an estimator
 - Well-calibrated probability intervals
- Both are useful!

The Bias-Variance Tradeoff

Reminder: an estimator is a random variable, an estimate is a constant

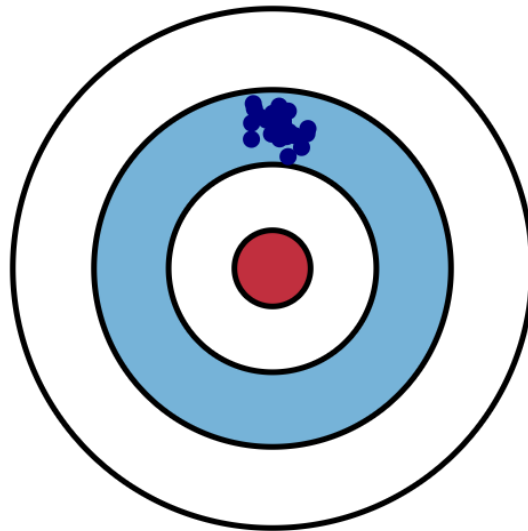
- *Bias*: systematic sampling error of the estimator
- *Variance*: variance of the estimator (from sampling & measurement error)
- Often we evaluate an estimator in terms of mean square error:
$$\text{MSE}(\hat{\theta}) = E_Y(\hat{\theta} - \theta)^2$$
- The Bias-Variance tradeoff: $\text{MSE}(\hat{\theta}) = \text{Var}(\hat{\theta}) + \text{Bias}(\hat{\theta})^2$

The Bias-Variance Tradeoff

- Variance of an estimator comes sampling from a population
 - If you were to repeatedly draw new samples of the same size how much would your estimates vary?
 - e.g. if $y_i \sim N(\mu, \sigma^2)$ then $\text{Var}(\bar{Y}) = \sigma^2/n$

Bias

The expected difference between the estimate and the response

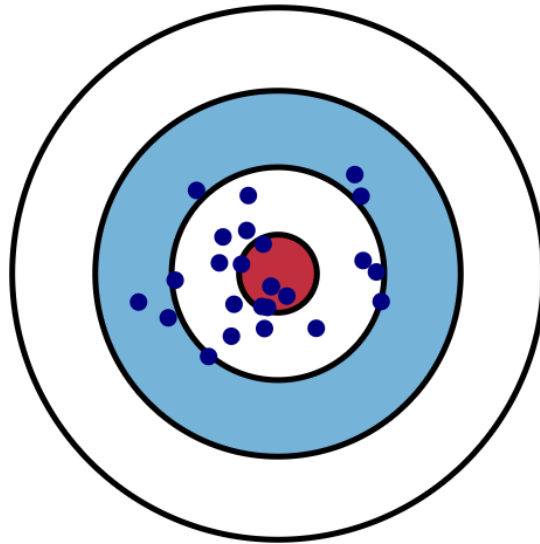


Statistical definition of bias:

$$E_Y[\hat{\theta} - \theta]$$

Variance

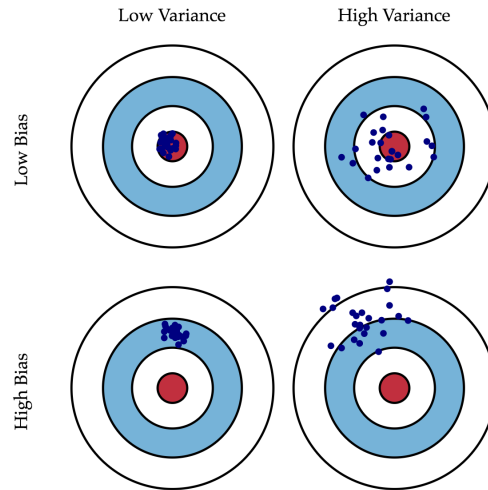
How variable is the prediction about its mean?



Statistical definition of variance:

$$E_Y[\hat{\theta} - E_Y[\hat{\theta}]]^2$$

Bias and Variance



$$\underbrace{\text{MSE}(\hat{\theta})}_{\text{accuracy}} = \underbrace{\text{Var}(\hat{\theta})}_{\text{variance}} + \underbrace{\left(E[\hat{\theta} - \theta] \right)^2}_{\text{bias squared}}$$

The Bias-Variance Tradeoff

- The prior distribution (usually) makes your estimator biased...
- But the prior distribution also (usually) reduces the variance!
- Example: compute the frequentist mean and variance of the posterior mean.

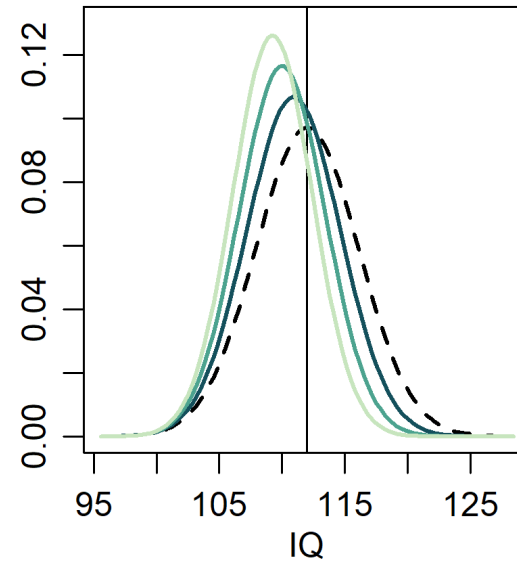
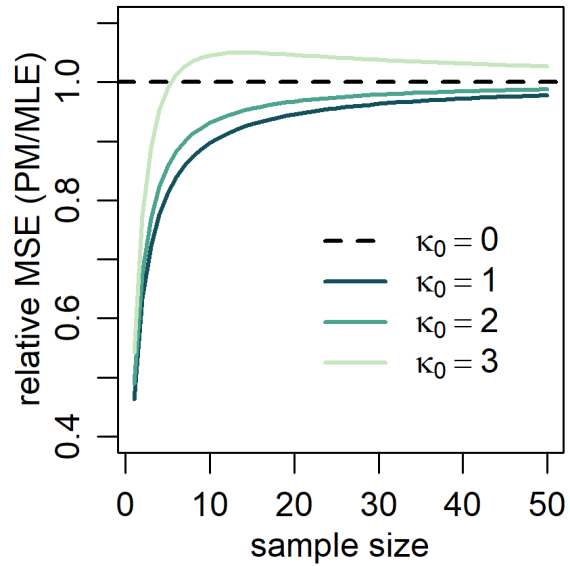
Example: IQ scores

- Scoring on **IQ tests** is designed to yield a $N(100, 15)$ distribution for the general population
- We observe IQ scores for a sample of n individuals from a particular town and estimate μ , the town-specific IQ score
- If we lacked knowledge about the town, a natural choice would be $\mu_0 = 100$
- Suppose the true parameters for this town are $\mu = 112$ and $\sigma = 13$
 - The town is smarter on average than the general population

Example: IQ scores

- What is the mean squared error of the MLE? MSE of the posterior mean?
- $\text{MSE}[\hat{\mu}_{MLE}] = \text{Var}[\hat{\mu}_{MLE}] = \frac{\sigma^2}{n} = \frac{169}{n}$
- $\text{MSE}[\hat{\mu}_{PM}|\theta_0] = w^2 \frac{169}{n} + (1 - w)^2 144$
- Reminder: $w = \frac{n}{\kappa_0 + n}$. For what values of n and κ_0 is the MSE smaller for the posterior mean estimator than the maximum likelihood?

Example: IQ scores



Decision Theory

Why the posterior mean?

- Often times we need to make a "decision" by providing a single estimate
- The posterior provides a full distribution over θ , which can be summarized in infinitely many ways
- Specify a *loss function* which describes the cost of estimating $\hat{\theta}$ when the truth is θ

Bayes Estimators

- The *loss function*: $L(\hat{\theta}, \theta)$
 - Squared error: $L(\hat{\theta}, \theta) = (\hat{\theta} - \theta)^2$
 - Absolute error: $L(\hat{\theta}, \theta) = |\hat{\theta} - \theta|$
- The **Bayes risk** is the posterior expected loss:

$$E_{\theta|y}[L(\hat{\theta}, \theta)] = \int L(\hat{\theta}, \theta)p(\theta | y)d\theta$$

- Choose an estimator of θ based on minimizing the Bayes risk.
- An estimator $\hat{\theta}$ is said to be a **Bayes estimator** if it minimizes the Bayes risk among all estimators.

Squared error loss

$$\min_{\hat{\theta}} E_{\theta|y}(\hat{\theta} - \theta)^2 = \min_{\hat{\theta}} \int (\hat{\theta} - \theta)^2 p(\theta | y) d\theta$$

Differentiate with respect to $\hat{\theta}$ and set equal to zero:

Absolute loss

$$\min_{\hat{\theta}} E_{\theta|y} |\hat{\theta} - \theta| = \min_{\hat{\theta}} \int |\hat{\theta} - \theta| p(\theta | y) d\theta$$

Differentiate with respect to $\hat{\theta}$ and set equal to zero:

Loss functions in practice

- Squared error and absolute error are good default loss functions
 - Motivated largely by mathematical considerations
- In practice we should define a loss function specific to our problem
- Loss in dollars? Loss in "quality of life"?

class: middle, center, inverse; background-image: none;

Normal distribution, unknown variance

Known mean, unknown variance

- Assume we have n observations with mean μ (known) and variance σ^2 (unknown)
- Define $d_i = (y_i - \mu)$ for notation convenience
- What is the likelihood $p(\sigma^2 \mid \mu, d_1, \dots, d_n)$?

Known mean, unknown variance

- If $X \sim \text{Gamma}(a, b)$, then $Y = \frac{1}{X} \sim \text{Inverse - Gamma}(a, b)$
- $p(y \mid a, b) = \frac{b^a}{\Gamma(a)} y^{-a-1} \exp\left\{-\frac{b}{y}\right\}$
- The inverse-gamma distribution is the conjugate prior distribution for the variance

Joint inference for normal parameters

- In practice, both μ and σ are unknown in the normal model.
- We've considered models when either μ is unknown but σ is known.
- In practice neither is known!
- Need to specify a joint prior distribution: $p(\mu, \sigma)$
- This is our first look at *multi-parameter* models.

Joint inference for the mean and variance

In the normal model we typically factorize the prior distribution $p(\mu, \sigma) = p(\mu | \sigma)p(\sigma)$.

Specifically:

$$\sigma^2 \propto \frac{1}{\sigma^2}$$

$$\mu | \sigma \sim \text{normal}(\mu_0, \sigma^2 / \kappa_0)$$

$$Y_1, \dots, Y_n | \mu, \sigma \sim \text{i.i.d. normal}(\mu, \sigma^2)$$

- μ_0 is interpreted as the prior sample mean
- κ_0 is a prior sample size

Example: midge wing length

- Modeling wing length of different species of midge (small, two-winged flies)
- From prior studies: mean wing length close to 1.9mm.
- Prior mean for μ is $\mu_0 = 1.9$ and non-informative prior for σ .
- Prior sample sizes: choose $\kappa_0 = 1$
- $(\bar{y}, s^2) = (1.804, 0.0169)$ are the sufficient statistics

Working with the log posterior

- As always, we will write down $p(\theta \mid y) \propto p(y \mid \theta)p(\theta)$
- In code, we always work with the **log-posterior** for numerical reasons
 - Mathematically it makes no difference, but computationally it is important
 - $L(\theta) \propto \prod p(y_i \mid \theta)$ is very small for moderate sample size (underflow)
 - $\ell(\theta) = \sum \log(p(y_i \mid \theta))$ is numerically stable
- Monte Carlo methods only require that we can evaluate the log posterior

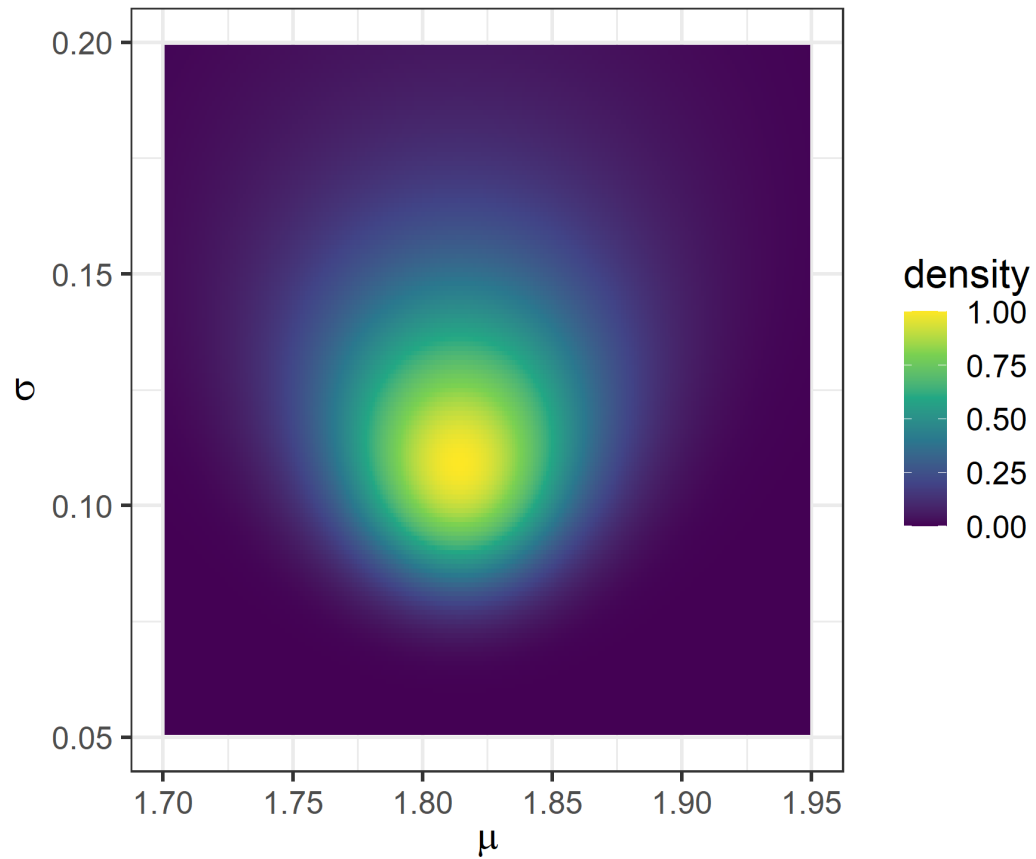
Grid approximation to the posterior distribution

```
log_normal_posterior <- Vectorize(function(mu, sigma) {  
  ### log likelihood  
  sum(dnorm(y, mu, sigma, log=TRUE)) +  
  ## plus log prior  
  dnorm(mu, mu0, sigma/sqrt(k0), log=TRUE) +  
  -2*log(sigma)  
})  
  
post_grid <- as_tibble(  
  expand.grid(seq(1.6, 2.0, by=0.001),  
              seq(0.01, 0.25, by=0.001)))  
colnames(post_grid) <- c("mu", "s")
```


Grid approximation to the posterior distribution

```
post_grid %>%  
  mutate(log_density = log_normal_posterior(mu, s)) %>%  
  mutate(density = exp(log_density - max(log_density))) %>%  
  ggplot() +  
  geom_raster(aes(mu, s, fill=exp(log_density))) +  
  xlim(c(1.7, 1.95)) + ylim(c(0.05, 0.2)) +  
  xlab(expression(mu)) +  
  ylab(expression(sigma)) +  
  theme_bw() +  
  scale_fill_continuous(type="viridis")
```

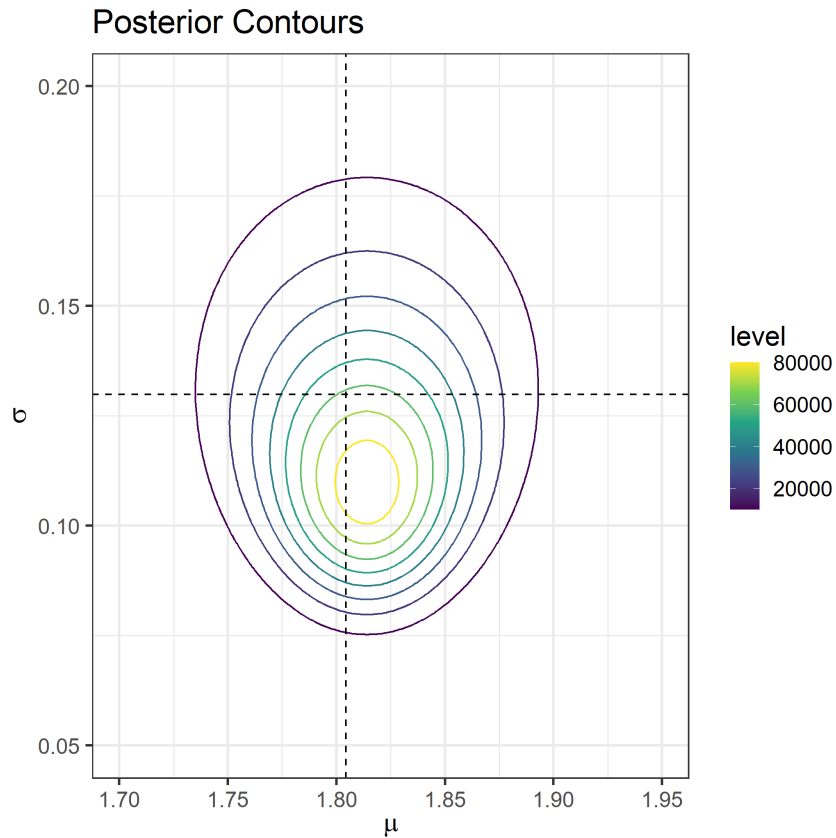
Grid approximation



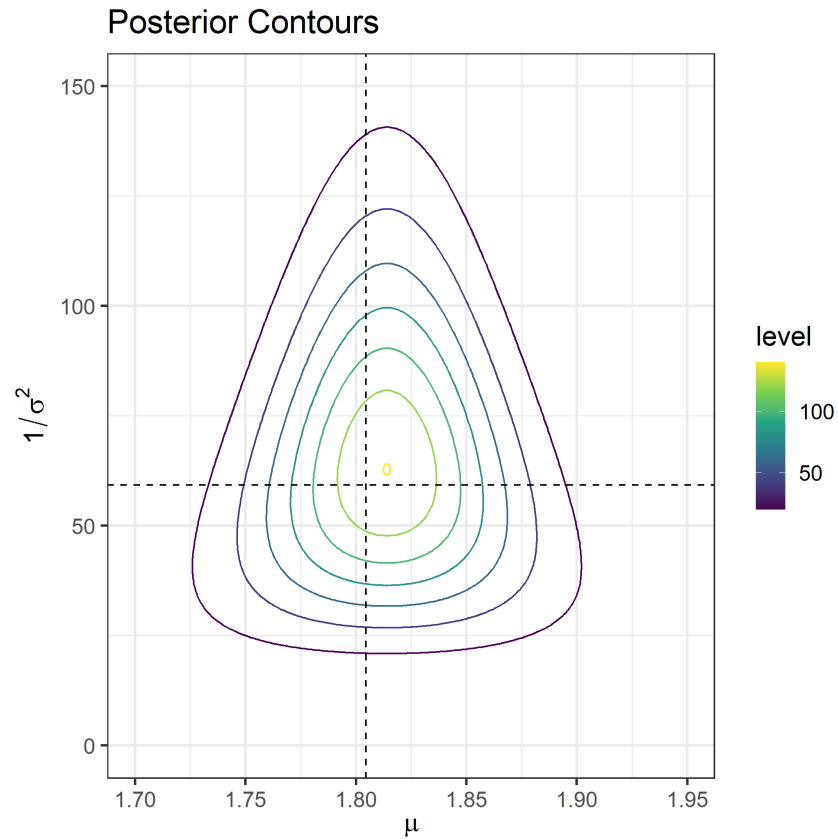
Contour Plot

```
post_grid %>%  
  mutate(density = exp(log_density - max(log_density))) %>%  
  ggplot() +  
  geom_contour(aes(mu, s, z=density, colour=stat(level)), size=2)  
  xlim(c(1.7, 1.95)) + ylim(c(0.05, 0.2)) +  
  xlab(expression(mu)) + ylab(expression(sigma)) +  
  ggtitle("Posterior Contours") +  
  theme_bw(base_size=16) +  
  scale_color_continuous(type="viridis")
```

Contour Plot (Standard Deviation)



Contour Plot (Precision)



Sampling from the joint posterior

- Contour and raster plots allow us to visualize the posterior (in two dimensions)
 - Need to know approximately where the high posterior density is (not easy)
- When we have more than 2 parameters visualization isn't feasible
- How do we summarize the posterior?
 - e.g. posterior means, posterior probabilities, intervals, etc..

Sampling from the joint posterior

- One approach: convert multi-parameter distribution into the product of many one parameter distributions.

Markov Chain Monte Carlo

- Markov Chain Monte Carlo (MCMC)
- More effective approach to sampling from multi-parameter distributions
- Samples in MCMC are **not** independent samples

MCMC Sampling with Stan

Stan



"A state-of-the-art platform for statistical modeling and high-performance statistical computation."

<http://mc-stan.org/>

Monte Carlo Sampling with Stan

- Stan is a "probabilistic programming language" with its own simple syntax
- R interface to Stan via `cmdstanr` package
- Define parameters θ and datatypes y
- Provide sampling model $p(y \mid \theta)$ and prior $p(\theta)$
- Stan translates model syntax into the log posterior density $\log(p(\theta \mid y))$ and automatically generates samples from this density

Example Stan File

```
// The input data is a vector 'y' of length 'N'.
data {
  int<lower=0> N;
  real<lower=0> k0;
  vector[N] y;
}

// The parameters accepted by the model. Our model
// accepts two parameters 'mu' and 'sigma'.
parameters {
  real mu;
  real<lower=0> sigma;
}

// The model to be estimated. We model the output
// 'y' to be normally distributed with mean 'mu'
// and standard deviation 'sigma'.
model {
  target += -2*log(sigma); //sigma prior
  mu ~ normal(1.9, sigma/k0); // mu prior
  y ~ normal(mu, sigma);
}
```

The Stan File

- A stan file ends in `.stan`
- Three important program blocks in a stan file:
 - Data block
 - Parameter block
 - Model block
 - Each block encapsulated in brackets, `{ ... }`.
- Stan needs data types:
 - `int`: integer valued data
 - `real`: continuous data or parameters
- Need to end every line with a semi-colon!
- Need to compile the Stan program before running.

Defining the input data in Stan

```
// The input data is a vector 'y' of length 'N'.  
data {  
  int<lower=0> N;  
  real<lower=0> k0;  
  vector[N] y;  
}
```

Defining the model parameters in Stan

```
// The parameters accepted by the model. Our model
// accepts two parameters 'mu' and 'sigma'.
parameters {
  real mu;
  real<lower=0> sigma;
}
```

Stan: The Basics

```
library(cmdstanr)

# compile stan model. This may take a minute.
stan_model <- cmdstan_model(stan_file="normal_model.stan")

## data is a list, arguments must match the
## arguments in data block of the stan file
stan_fit <- stan_model$sample(data=list(N=n, y=y, k0=0.1),
                             refresh=0)

## Running MCMC with 4 sequential chains...
##
## Chain 1 finished in 0.0 seconds.
## Chain 2 finished in 0.0 seconds.
## Chain 3 finished in 0.0 seconds.
## Chain 4 finished in 0.0 seconds.
##
## All 4 chains finished successfully.
## Mean chain execution time: 0.0 seconds.
## Total execution time: 0.8 seconds.
```

Stan: The Basics

```
## Convert Stan output to a list of MC samples
samples <- stan_fit$draws(format="df")
samples
```

```
## # A draws_df: 1000 iterations, 4 chains, and 3 variables
##   lp__      mu sigma
## 1      17  1.8 0.149
## 2      16  1.8 0.163
## 3      16  1.8 0.095
## 4      17  1.8 0.124
## 5      17  1.8 0.109
## 6      13  1.7 0.078
## 7      14  1.9 0.142
## 8      14  1.9 0.142
## 9      15  1.9 0.153
## 10     16  1.9 0.096
## # ... with 3990 more draws
## # ... hidden reserved variables {'.chain', '.iteration', '.draw'}
```


Stan: The Basics

```
## names in list are the defined parameters
names(samples)
```

```
## [1] "lp__"      "mu"        "sigma"     ".chain"    ".iteration"
## [6] ".draw"
```

```
mu_samples <- samples$mu
sigma_samples <- samples$sigma
```

```
## By default there will be 4000 samples
length(mu_samples)
```

```
## [1] 4000
```

Stan: The Basics

```
## Information about MC samples  
summary(mu_samples)
```

```
##      Min. 1st Qu.  Median      Mean 3rd Qu.      Max.  
##  1.676   1.781   1.807   1.808   1.833   2.054
```

```
summary(sigma_samples)
```

```
##      Min. 1st Qu.  Median      Mean 3rd Qu.      Max.  
## 0.06275 0.10294 0.11922 0.12532 0.14163 0.40957
```

```
summary(1/sigma_samples^2)
```

```
##      Min. 1st Qu.  Median      Mean 3rd Qu.      Max.  
##  5.961  49.853  70.360  75.399  94.375 253.983
```

Stan: The Basics

Shortcut for summary statistics:

```
## Information about MC samples
stan_fit$summary()
```



```
## # A tibble: 3 × 10
##   variable    mean median      sd    mad      q5     q95   rhat ess_bulk ess_
##   <chr>      <dbl> <dbl>   <dbl> <dbl>   <dbl> <dbl> <dbl>   <dbl>
## 1 lp__      16.4   16.8   1.09   0.766  14.3   17.5   1.00   1277.
## 2 mu        1.81   1.81   0.0416 0.0393  1.74   1.87   1.00   2055.
## 3 sigma     0.125  0.119  0.0329 0.0281  0.0845 0.186  1.00   2252.
```

Visualizing Posterior Samples from Stan

```
tibble(Mean = mu_samples, Precision=1/sigma_samples^2) %>%  
  ggplot() +  
  geom_point(aes(x=Mean, y=Precision)) +  
  theme_bw(base_size=16)
```

