
author: "Rory Quinlan" output: html_document —

```
# Descriptive Stats
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n_a = length(Y_a)
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n_b = length(Y_b)
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ybar_a = mean(Y_a)
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```
ybar_b = mean(Y_b)
```

```
# Set param values
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```
a_theta = 2
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```
b_theta = 1
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```
S = 5000
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```
ab_gamma = c(8, 16, 32, 64, 128)
```

```

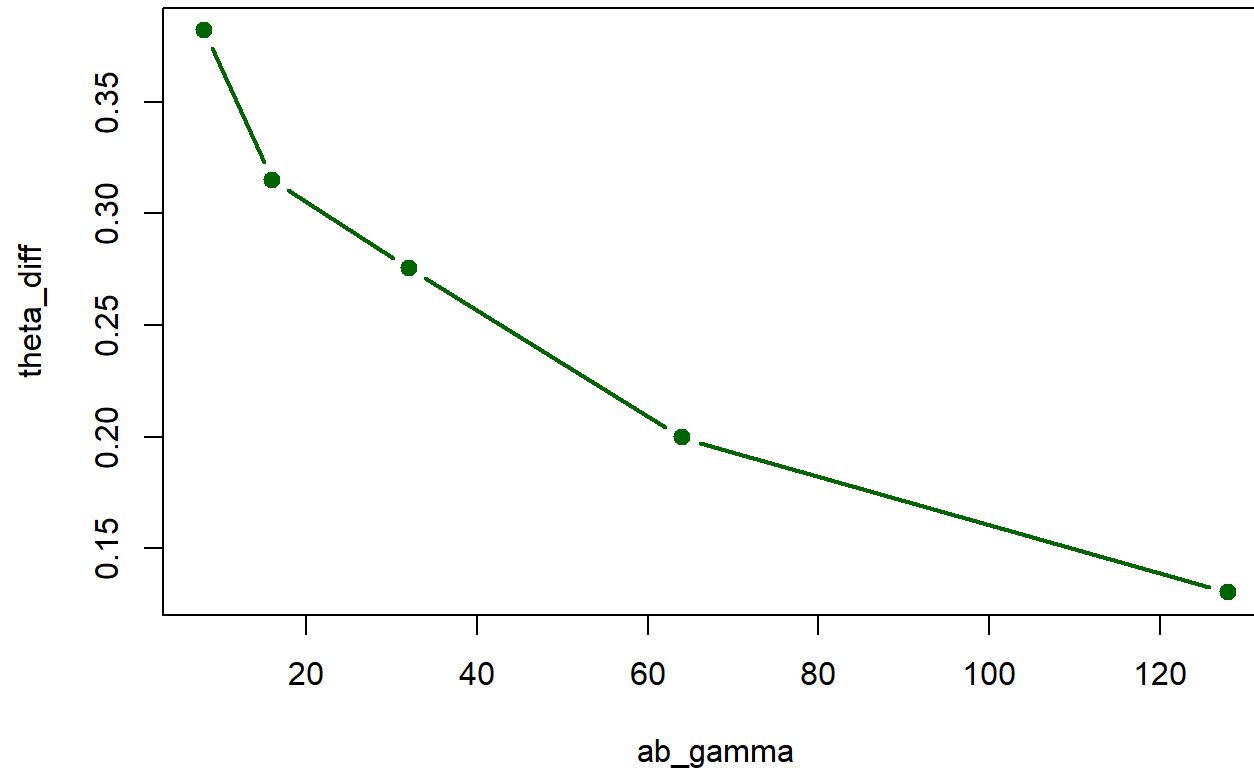
theta_diff = sapply(ab_gamma, function(abg) {
  a_gamma = b_gamma = abg
  THETA = numeric(S)
  GAMMA = numeric(S)
  # Starting values
  theta = ybar_a
  # Relative rate theta_B /theta_A
  gamma = ybar_a / ybar_b
  # For each value in s (5,000) create a theta with gamma(1,...)
  # Then create a a gamma (1,...)
  for (s in 1:S) {
    theta = rgamma(
      1,
      a_theta + n_a * ybar_a + n_b * ybar_b,
      b_theta + n_a + n_b * gamma
    )
    gamma = rgamma(
      1,
      a_gamma + n_b * ybar_b,
      b_gamma + n_b * theta
    )
    THETA[s] = theta
    GAMMA[s] = gamma
  }
  # Reconstruct theta_ and theta_B
  THETA_A = THETA
  THETA_B = THETA * GAMMA
  mean(THETA_B - THETA_A)
})

```

```

# Plot
plot(x = ab_gamma, y = theta_diff, xlab = "ab_gamma", ylab = "theta_diff", pch=19, lwd=2, col="darkgreen", type="b")

```



- Since γ_a and γ_b are equal, the gamma distribution will be centered at 1. As our belief in that increases, the mean posterior difference between θ_B and θ_A decreases.