

# Non-Conjugate Models and Grid Approximations

Patrick Lam

# Outline

The Binomial Model with a Non-Conjugate Prior

Bayesian Regression with Grid Approximations

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> triangle.prior <- function(x) {  
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+     8 * x  
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To find the **posterior**, we need to use grid approximation methods.

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There are two ways to get the **normalized posterior** in R.

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2. Evaluate each grid point  $(\pi_0 + k), (\pi_0 + 2k), \dots, (\pi_0 + mk)$  at the **unnormalized posterior**:  $p(y|\pi_0 + ik)p(\pi_0 + ik)$ .



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```
> unnormal.post.ord <- posterior.function(theta = grid.points,  
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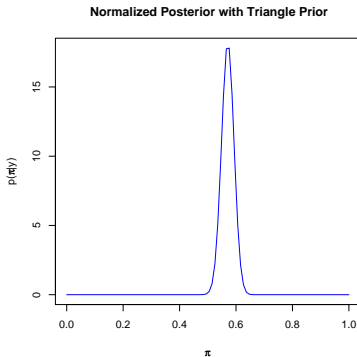
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```
> post.ord <- unnormal.post.ord/normal.constant
```

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```
> plot(x = grid.points, y = post.ord, type = "l", col = "blue",  
+       xlab = expression(pi), ylab = expression(paste("p(", pi,  
+       "|y)")), main = "Normalized Posterior with Triangle Prior")
```





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```
> set.seed(12345)  
> posterior.triangle.2 <- sample(grid.points, size = 10000, replace = T,  
+   prob = unnormal.post.ord)  
> all.equal(posterior.triangle.1, posterior.triangle.2)
```

```
[1] TRUE
```

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Let's use a couple of Normal priors on  $\alpha$  and  $\beta$ .

$$\text{Posterior}(\alpha, \beta) \propto \prod_{i=1}^n \text{Poisson}(\lambda_i) \times \text{Normal}(\mu_\alpha, \sigma_\alpha^2) \times \text{Normal}(\mu_\beta, \sigma_\beta^2)$$

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For the Poisson, the link function is the natural log function:

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However, typically we use the **inverse link function** to reparameterize  $\alpha + \beta x_i$  so that it can only take on positive values.

$$\lambda_i = \exp(\alpha + \beta x_i)$$



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Probit:

- ▶ Link:  $\Phi^{-1}(\pi_i) = \alpha + \beta x_i$
- ▶ Inverse Link:  $\pi_i = \Phi(\alpha + \beta x_i)$

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So our posterior is

$$p(\alpha, \beta | \mathbf{y}, \mathbf{x}) \propto \prod_{i=1}^n \frac{e^{-\lambda_i} \lambda_i^{y_i}}{y_i!} \times \frac{1}{\sqrt{2\pi\sigma_\alpha^2}} \exp\left(-\frac{(\alpha - \mu_\alpha)^2}{2\sigma_\alpha^2}\right) \\ \times \frac{1}{\sqrt{2\pi\sigma_\beta^2}} \exp\left(-\frac{(\beta - \mu_\beta)^2}{2\sigma_\beta^2}\right)$$

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Using our inverse link function, we get

$$p(\alpha, \beta | \mathbf{y}, \mathbf{x}) \propto \prod_{i=1}^n \frac{e^{-\exp(\alpha + \beta x_i)} (\exp(\alpha + \beta x_i))^{y_i}}{y_i!} \times \frac{1}{\sqrt{2\pi\sigma_\alpha^2}} \exp\left(-\frac{(\alpha - \mu_\alpha)^2}{2\sigma_\alpha^2}\right) \\ \times \frac{1}{\sqrt{2\pi\sigma_\beta^2}} \exp\left(-\frac{(\beta - \mu_\beta)^2}{2\sigma_\beta^2}\right)$$

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We can combine and simplify a bunch of terms ... or we can use some canned R functions.

$$\ln \text{Posterior}(\alpha, \beta) \propto \sum_{i=1}^n \ln \text{Poisson}(\exp(\alpha + \beta \mathbf{x}_i)) + \ln \text{Normal}(\mu_\alpha, \sigma_\alpha^2) \\ + \ln \text{Normal}(\mu_\beta, \sigma_\beta^2)$$

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```
> poisson.posterior <- function(theta, y, x, prior.mean.a, prior.var.a,
+   prior.mean.b, prior.var.b) {
+   a <- theta[1]
+   b <- theta[2]
+   lambda <- exp(a + b * x)
+   log.like <- sum(dpois(y, lambda = lambda, log = T))
+   log.prior.a <- dnorm(a, mean = prior.mean.a, sd = sqrt(prior.var.a),
+     log = T)
+   log.prior.b <- dnorm(b, mean = prior.mean.b, sd = sqrt(prior.var.b),
+     log = T)
+   log.post <- log.like + log.prior.a + log.prior.b
+   return(exp(log.post))
+ }
```

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```
> mu.a <- mu.b <- 0
> sigma2.a <- sigma2.b <- 20
```

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```
> mle <- glm(num ~ coop, data = sanction, family = poisson)$coef  
> mle.se <- summary(glm(num ~ coop, data = sanction,  
+   family = poisson))$coef[, 2]
```

I then create a grid that is centered at the MLE and spans an arbitrary 5 standard errors away on both sides.

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```
> grid.a <- seq(from = mle[1] - 5 * mle.se[1], to = mle[1] + 5 *  
+   mle.se[1], length.out = 200)  
> grid.b <- seq(from = mle[2] - 5 * mle.se[2], to = mle[2] + 5 *  
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> post.ord <- apply(grid.points, MARGIN = 1, FUN = poisson.posterior,  
+   y = y.vec, x = x.vec, prior.mean.a = mu.a, prior.var.a = sigma2.a,  
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+   prior.mean.b = mu.b, prior.var.b = sigma2.b)
```

and then sample (letting R normalize for us) to get the simulated posterior.

```
> sample.indices <- sample(1:nrow(grid.points), size = 10000, replace = T,  
+   prob = post.ord)  
> sim.posterior <- grid.points[sample.indices, ]
```

# Posterior Summary

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      Var1    Var2
-1.007    1.207
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```
> posterior.sd <- apply(sim.posterior, MARGIN = 2, FUN = sd)
> posterior.sd
```

```
      Var1    Var2
0.1466    0.0453
```

```
> mle.se
```

```
(Intercept)      coop
      0.14602      0.04504
```