Estimating a Gamma distribution

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Abstract

This note derives a fast algorithm for maximum-likelihood estimation of both parameters of a Gamma distribution or negative-binomial distribution.

1 Introduction

We have observed n independent data points $X = [x_1..x_n]$ from the same density θ . We restrict θ to the class of Gamma densities, i.e. $\theta = (a, b)$:

$$p(x|a,b) = \operatorname{Ga}(x;a,b) = \frac{x^{a-1}}{\Gamma(a)b^a} \exp(-\frac{x}{b})$$

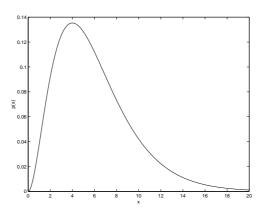


Figure 1: The Ga(3,2) density function.

Figure 1 plots a typical Gamma density. In general, the mean is ab and the mode is (a-1)b.

2 Maximum likelihood

The log-likelihood is

$$\log p(D|a,b) = (a-1)\sum_{i} \log x_i - n\log \Gamma(a) - na\log b - \frac{1}{b}\sum_{i} x_i$$
 (1)

$$= n(a-1)\overline{\log x} - n\log\Gamma(a) - na\log b - n\bar{x}/b$$
 (2)

The maximum for b is easily found to be

$$\hat{b} = \bar{x}/a \tag{3}$$

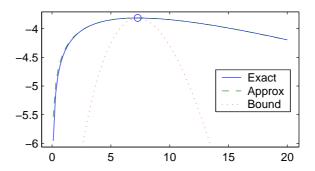


Figure 2: The log-likelihood (4) versus the Gamma-type approximation (9) and the bound (6) at convergence. The approximation is nearly identical to the true likelihood. The dataset was 100 points sampled from Ga(7.3, 4.5).

Substituting this into (1) gives

$$\log p(D|a, \hat{b}) = n(a-1)\overline{\log x} - n\log\Gamma(a) - na\log\bar{x} + na\log a - na$$
(4)

We will describe two algorithms for maximizing this function.

The first method will iteratively maximize a lower bound. Because $a \log a$ is convex, we can use a linear lower bound:

$$a \log a \ge (1 + \log a_0)(a - a_0) + a_0 \log a_0$$
 (5)

$$\log p(D|a,\hat{b}) \geq n(a-1)\overline{\log x} - n\log\Gamma(a) - na\log\bar{x} + n(1+\log a_0)(a-a_0) + na_0\log a_0 - na$$
 (6)

The maximum is at

$$0 = n\overline{\log x} - n\Psi(a) - n\log \bar{x} + n(1 + \log a_0) - n$$

$$\Psi(\hat{a}) = \overline{\log x} - \log \bar{x} + \log a_0$$
(8)

$$\Psi(\hat{a}) = \overline{\log x} - \log \bar{x} + \log a_0 \tag{8}$$

where Ψ is the digamma function. The iteration proceeds by setting a_0 to the current \hat{a} , then inverting the Ψ function to get a new \hat{a} . Because the log-likelihood is concave, this iteration must converge to the (unique) global maximum. Unfortunately, it can be quite slow, requiring around 250 iterations if a = 10, less for smaller a, and more for larger a.

The second algorithm is much faster, and is obtained via 'generalized Newton' [1]. Using an approximation of the form,

$$\log p(D|a,\hat{b}) \approx c_0 + c_1 a + c_2 \log(a) \tag{9}$$

the update is

$$\frac{1}{a^{new}} = \frac{1}{a} + \frac{\overline{\log x} - \log \bar{x} + \log a - \Psi(a)}{a^2 (1/a - \Psi'(a))}$$
(10)

This converges in about four iterations. Figure 2 shows that this approximation is very close to the true loglikelihood, which explains the good performance.

A good starting point for the iteration is obtained via the approximation

$$\log \Gamma(a) \approx a \log(a) - a - \frac{1}{2} \log a + \text{const.}$$
 (Stirling) (11)

$$\Psi(a) \approx \log(a) - \frac{1}{2a}$$

$$\hat{a} \approx \frac{0.5}{\log \overline{x} - \overline{\log x}}$$

$$\tag{12}$$

$$\hat{a} \approx \frac{0.5}{\log \overline{x} - \overline{\log x}} \tag{13}$$

(Note that $\log \overline{x} \ge \overline{\log x}$ by Jensen's inequality.)

Negative binomial 2.1

The maximum-likelihood problem for the negative binomial distribution is quite similar to that for the Gamma. This is because the negative binomial is a mixture of Poissons, with Gamma mixing distribution:

$$p(x|a,b) = \int_{\lambda} \text{Po}(x;\lambda) \text{Ga}(\lambda;a,b) d\lambda = \int_{\lambda} \frac{\lambda^{x}}{x!} e^{-\lambda} \frac{\lambda^{a-1}}{\Gamma(a)b^{a}} e^{-\lambda/b} d\lambda$$
 (14)

$$= \left(\begin{array}{c} a+x-1\\ x \end{array}\right) \left(\frac{b}{b+1}\right)^x \left(1-\frac{b}{b+1}\right)^a \tag{15}$$

Let's consider a slightly generalized negative binomial, where the 'waiting time' for x is given by t:

$$p(x|t,a,b) = \int_{\lambda} \text{Po}(x;\lambda t) \text{Ga}(\lambda;a,b) d\lambda = \int_{\lambda} \frac{(\lambda t)^{x}}{x!} e^{-\lambda t} \frac{\lambda^{a-1}}{\Gamma(a)b^{a}} e^{-\lambda/b} d\lambda$$
 (16)

$$= \left(\begin{array}{c} a+x-1\\ x \end{array}\right) \left(\frac{bt}{bt+1}\right)^x \left(1-\frac{bt}{bt+1}\right)^a \tag{17}$$

Given a data set $D = \{(x_i, t_i)\}$, we want to estimate (a, b). One approach is to use EM, where the E-step infers the hidden variable λ_i :

$$E[\lambda_i] = (x_i + a) \frac{b}{bt_i + 1} \tag{18}$$

$$E[\log \lambda_i] = \Psi(x_i + a) + \log \frac{b}{bt_i + 1}$$
(19)

The M-step then maximizes

$$(a-1)\sum_{i} E[\log \lambda_{i}] - n\log \Gamma(a) - na\log b - \frac{1}{b}\sum_{i} E[\lambda_{i}]$$
(20)

which is a Gamma maximum-likelihood problem.

References

[1] Thomas P. Minka. Beyond newton's method. research.microsoft.com/~minka/papers/newton.html,