${\sf Andrew\ Gard\ -\ equitable.equations@gmail.com}$



The Gamma Distribution in R



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Base R functions use the rate parameter λ by default.





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Here and elsewhere, the scale parameter θ may be used instead, but it must be named explicitly.

```
rgamma(6, 2, scale = 2)
## [1] 6.891120 2.254109 1.302274 4.101624 7.770818 5.081080
```



• pgamma(x, alpha, lambda) is the cumulative distribution function. It returns the probability that the waiting time for alpha occurrences is no more than x. As usual in R, x can be a vector.



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qgamma(p, alpha, lambda) is the inverse distribution function. It returns
the value x such that pgamma(x, alpha, lambda) = p. In other words, it
computes quantiles in the specified gamma distribution. Again, p can be a
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the value x such that pgamma(x, alpha, lambda) = p. In other words, it
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vector. For instance,

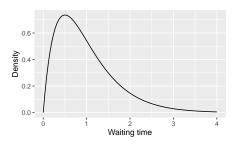
```
qgamma(c(.2, .4, .6, .8), 2, .5)
## [1] 1.648777 2.752843 4.044626 5.988617
```



• dgamma(x, alpha, lambda) is the probability density function (pdf) of the gamma distribution with shape parameter alpha and rate parameter lambda.



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Example. Calls to a customer-service line come at an average rate of 1 every 3 minutes.

1. What is the probability that more than an hour elapses before 25 calls come in?

2. What is the 95^{th} percentile for time needed for 5 calls to come in?

3. Simulate waiting times for 5 calls 1000 times. Plot the results.

