Common Probability Distributions

R Function Conventions

R provides functions to do four different types of things with a probability distribution:

Task	Function name starts with
Evaluate the density function $f(x)$	d
Calculate the p robability $P(X \leq x)$	p
Find a quantile	q
Generate a \mathbf{r} andom number	r

I'll illustrate each of these with the normal distribution:

1. Evaluate f(x), the density function (pdf for continuous random variables or pmf for discrete random variables)

```
dnorm(0.5, mean = 0, sd = 1, log = FALSE)

## [1] 0.3520653

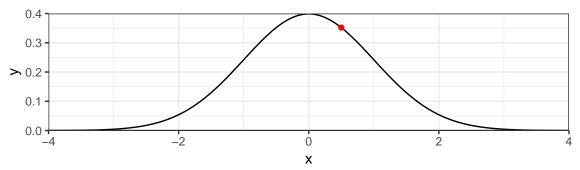
This gives the vertical axis coordinate of the pdf (or pmf) evaluated at x = 0.5:

# evaluate the normal(0, 1) pdf at x = 0.5 and store in a data frame
evaluated_density_to_plot <- data.frame(
    x = 0.5,
    y = dnorm(0.5, mean = 0, sd = 1, log = FALSE)</pre>
```

```
## x y
## 1 0.5 0.3520653
```

evaluated_density_to_plot

```
# make a plot of the density function with a red point at the (x, y) pair found above
library(ggplot2)
ggplot(data = data.frame(x = c(-4, 4)), mapping = aes(x = x)) +
    stat_function(fun = dnorm) +
    geom_point(data = evaluated_density_to_plot, mapping = aes(x = x, y = y), color = "red") +
    coord_cartesian(xlim = c(-4, 4), ylim = c(0, 0.4), expand = FALSE) +
    theme_bw()
```



2. Calculate $P(X \leq q)$, the probability that a value drawn from this distribution is less than q

```
pnorm(0.5, mean = 0, sd = 1, log = FALSE)
```

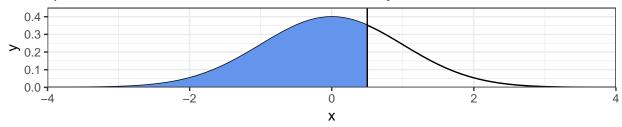
[1] 0.6914625

This calculates $P(X \le 0.5)$ if $X \sim \text{Normal}(0,1)$, the shaded area below:

```
x_grid <- seq(from = -4, to = 0.5, length = 101)
region_to_shade <- data.frame(
    x = c(-4, x_grid, 0.5),
    y = c(0, dnorm(x_grid, mean = 0, sd = 1, log = FALSE), 0)
)

ggplot(data = data.frame(x = c(-4, 4)), mapping = aes(x = x)) +
    stat_function(fun = dnorm) +
    geom_polygon(
    mapping = aes(x = x, y = y),
    fill = "cornflowerblue",
    data = region_to_shade) +
    geom_vline(xintercept = 0.5) +
    coord_cartesian(xlim = c(-4, 4), ylim = c(0, 0.45), expand = FALSE) +
    theme_bw() +
    ggtitle("pnorm: find the shaded area if you know the location of the vertical line\nqnorm: find the location</pre>
```

pnorm: find the shaded area if you know the location of the vertical line qnorm: find the location of the vertical line if you know the shaded area



3. Find quantiles: For a given number p, find the value q such that $P(X \le q) = p$

```
qnorm(0.6914625, mean = 0, sd = 1, log = FALSE)
```

[1] 0.5000001

This calculates the location of the vertical line in the plot above, given that the area to the left of that line is 0.6914625: $P(X \le ?) = 0.6914625$

4. Generate random numbers

```
rnorm(n = 5, mean = 0, sd = 1)
```

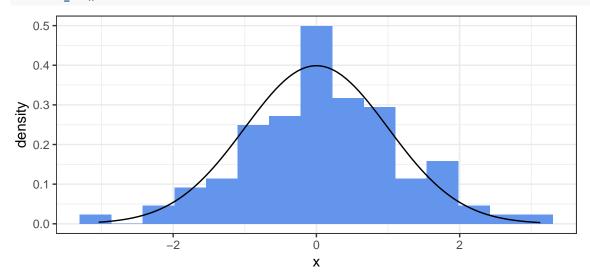
```
## [1] -1.6494928 -2.3563596  0.6839980 -0.5959463  0.8583114
```

If we want to plot these values using ggplot2, it's convenient to store them as a variable in a data frame:

```
# set a seed for random number generation so that I get reproducible results
set.seed(59749)

# create a new data frame called data_to_plot, with one variable in it: x
data_to_plot <- data.frame(
    x = rnorm(n = 100, mean = 0, sd = 1)
)

# make a plot
ggplot(data = data_to_plot, mapping = aes(x = x)) +
    geom_histogram(mapping = aes(y = ..density..),
    fill = "cornflowerblue",
    bins = 15) +
    stat_function(fun = dnorm) +
    theme_bw()</pre>
```



Discrete Distributions

Bernoulli(p)

X = the result of a single experiment with one of two outcomes ("success", coded as 1, or "failure", coded as 0), where the probability of success is p.

```
parameters p = \text{probability of success}

p.f. f(x|p) = p^x (1-p)^{(1-x)}

Mean p

Variance p(1-p)

R functions dbinom(..., size = 1, prob = p), pbinom, qbinom, rbinom
```

Binomial(n, p)

X = the total number of successes in n independent and identically distributed Bernoulli trials, each with probability of success p.

```
parameters n= number of trials, p= probability of success p.f. f(x|n,p)=\binom{n}{x}p^x(1-p)^{(n-x)} Mean np Variance np(1-p) R functions dbinom(..., size = n, prob = p), pbinom, qbinom, rbinom
```

Uniform(a, b)

X = an integer between a and b (inclusive), where each integer from a to b is equally likely.

```
parameters a: lower endpoint, b: upper endpoint  \begin{array}{ll} p.f. & f(x|a,b) = \frac{1}{b-a+1} \\ Mean & \frac{a+b}{2} \\ Variance & \frac{(b-a)(b-a+2)}{12} \\ R \text{ functions} & No \text{ specific d, p, or q functions. For random number generation, you could use } \\ & \text{sample(seq(from = a, to = b, by = 1), size = n, replace = TRUE)} \end{array}
```

Geometric(p)

X = the number of failures that occur before the first success in a sequence of independent and identically distributed Bernoulli trials.

```
parameters p \in (0,1): probability of success on each trial p.f. f(x|p) = p(1-p)^x Mean \frac{1-p}{p} Variance \frac{1-p}{p^2} R functions p \in (0,1): prob = p, ...), pgeom, qgeom, rgeom
```

Be careful – there are other parameterizations used in other sources.

Negative Binomial(r, p)

X = the number of failures which occur in a sequence of independent and identically distributed Bernoulli trials before r successes occur.

```
parameters r>0: target number of successes, p>0 = probability of success on each trial p.f. f(x|r,p) = {r+x-1 \choose x} p^r (1-p)^x
Mean \frac{r(1-p)}{p}
Variance \frac{r(1-p)}{p^2}
R functions dnbinom(..., size = r, prob = p, ...), pnbinom, qnbinom, rnbinom
```

Be careful – there are multiple other parameterizations used in other sources.

Hypergeometric (A, B, n)

X = the number of successes in n draws (without replacement) from a finite population that contains exactly A successes and B failures.

```
parameters A: number of successes in the population, B: number of failures in the population, n: sample size  \begin{array}{ll} \text{p.f.} & f(x|A,B,n) = \frac{\binom{A}{x}\binom{B}{n-x}}{\binom{A+B}{n}} \\ \text{Mean} & \frac{nA}{A+B} \\ \text{Variance} & \frac{nAB}{(A+B)^2} \frac{A+B-n}{A+B-1} \\ \text{R functions} & \text{dhyper(..., m = A, n = B, k = n, ...), phyper, qhyper, rhyper} \end{array}
```

$Poisson(\lambda)$

X = the number of events occurring in a fixed interval of time or space if these events occur with a known constant rate and independently of the time since the last event.

```
parameters \lambda: rate parameter p.f. f(x|\lambda) = e^{-\lambda} \frac{\lambda^x}{x!} Mean \lambda Variance \lambda R functions dpois(..., lambda = lambda), ppois, qpois, rpois
```

Multinomial(n, p), where $p = (p_1, p_2, \dots, p_k)$

 $X = (X_1, X_2, ..., X_k)$ = the vector of counts for how many observations fell into each of k categories in a sample of n independent trials where the item sampled in each trial falls into category j with probability p_j . (Roll a weighted die with k sides n times. How many times did each face of the die come up?)

```
parameters n: number of trials, p=(p_1,p_2,\ldots,p_k): vector of probabilities for each category p.f. f(x|\lambda) = \frac{n!}{x_1!x_2!\cdots x_k!}p_1^{x_1}p_2^{x_2}\cdots p_k^{x_k}
Mean E(X_i) = np_i
Variance Var(X_i) = np_i(1-p_i)
R functions dmultinom(..., size = n, prob = p), rmultinom
```

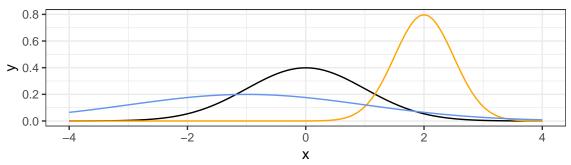
Continous Distributions

$Normal(\mu, \sigma^2)$

A real number.

```
parameters \mu: mean, \sigma^2 > 0: variance  \begin{array}{ll} \text{p.f.} & f(x|\mu,\sigma^2) = (2\pi\sigma^2)^{-\frac{1}{2}}e^{-\frac{1}{2}\frac{(x-\mu)^2}{\sigma^2}} \\ \text{Mean} & \mu \\ \text{Variance} & \sigma^2 \\ \text{R functions} & \texttt{dnorm(..., mean = }\mu\text{, sd = }\sigma\text{), pnorm, qnorm, rnorm} \end{array}
```

```
ggplot(data = data.frame(x = c(-4, 4)), mapping = aes(x = x)) +
   stat_function(fun = dnorm, args = list(mean = 0, sd = 1), color = "black") +
   stat_function(fun = dnorm, args = list(mean = 2, sd = 0.5), color = "orange") +
   stat_function(fun = dnorm, args = list(mean = -1, sd = 2), color = "cornflowerblue") +
   theme_bw()
```

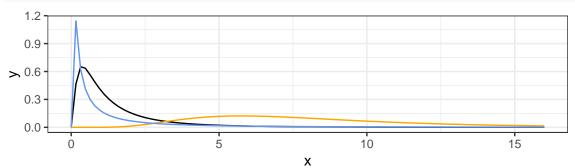


$\mathbf{Lognormal}(\mu, \sigma^2)$

A positive number. If $X \sim \text{Lognormal}(\mu, \sigma^2)$ and $Y = \log(X)$, then $Y \sim \text{Normal}(\mu, \sigma^2)$.

```
parameters \mu: mean of \log(X), \sigma^2 > 0: variance of \log(X) p.f. f(x|\mu,\sigma^2) = x^{-1}(2\pi\sigma^2)^{-\frac{1}{2}} \exp\left[-\frac{1}{2}\frac{\{\log(x)-\mu\}^2}{\sigma^2}\right] Mean \exp\left(\mu + \frac{\sigma^2}{2}\right) Variance \{\exp(\sigma^2) + 1\} \exp(2\mu + \sigma^2) R functions dlnorm(..., meanlog = \mu, sdlog = \sigma), plnorm, qlnorm, rlnorm
```

```
ggplot(data = data.frame(x = c(0, 16)), mapping = aes(x = x)) +
   stat_function(fun = dlnorm, args = list(meanlog = 0, sdlog = 1), color = "black") +
   stat_function(fun = dlnorm, args = list(meanlog = 2, sdlog = 0.5), color = "orange") +
   stat_function(fun = dlnorm, args = list(meanlog = -1, sdlog = 2), color = "cornflowerblue") +
   theme_bw()
```



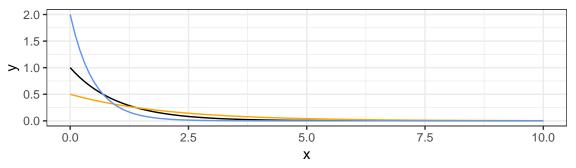
Exponential(λ)

A non-negative real number. Often used as a model for waiting times – but need to check whether this is a good model for a given data set.

```
parameters \lambda > 0: rate parameter p.f. f(x|\lambda) = \lambda e^{-\lambda x}
Mean \frac{1}{\lambda}
Variance \frac{1}{\lambda^2}
R functions dexp(..., rate = \lambda), pexp, qexp, rexp
```

Be careful – in some sources, the exponential distribution is parameterized in terms of a scale parameter $\beta = \frac{1}{\lambda}$

```
ggplot(data = data.frame(x = c(0, 10)), mapping = aes(x = x)) +
   stat_function(fun = dexp, args = list(rate = 1), color = "black") +
   stat_function(fun = dexp, args = list(rate = 0.5), color = "orange") +
   stat_function(fun = dexp, args = list(rate = 2), color = "cornflowerblue") +
   theme_bw()
```



$Gamma(\alpha, \lambda)$

A non-negative real number. Often used as a model for waiting times – but need to check whether this is a good model for a given data set.

```
parameters \alpha \geq 0: shape parameter, \lambda > 0: rate parameter p.f. f(x|\alpha,\lambda) = \frac{\lambda^{\alpha}}{\Gamma(\alpha)}x^{\alpha-1}e^{-\lambda x}

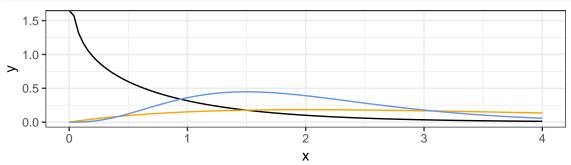
Mean \frac{\alpha}{\lambda}

Variance \alpha\lambda^2

R functions dgamma(..., shape = \alpha, rate = \lambda), pgamma, qgamma, rgamma
```

Be careful – there are multiple other parameterizations used in other sources.

```
ggplot(data = data.frame(x = c(0, 4)), mapping = aes(x = x)) +
   stat_function(fun = dgamma, args = list(shape = 0.8, rate = 1), color = "black") +
   stat_function(fun = dgamma, args = list(shape = 2, rate = 0.5), color = "orange") +
   stat_function(fun = dgamma, args = list(shape = 4, rate = 2), color = "cornflowerblue") +
   theme_bw()
```

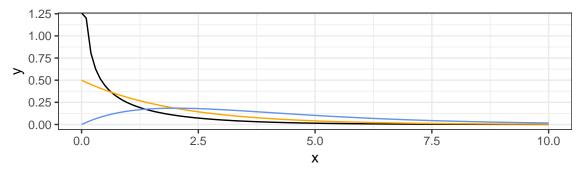


Chi-Squared(n)

A non-negative real number.

```
parameters n: degrees of freedom  \text{p.f.} \quad f(x|n) = \frac{1}{2^{n/2}\Gamma(n/2)} x^{\frac{n}{2}-1} e^{-\frac{x}{2}}  Mean n Variance 2n R functions \text{dchisq(..., df = n), pchisq, qchisq, rchisq}
```

```
ggplot(data = data.frame(x = c(0, 10)), mapping = aes(x = x)) +
stat_function(fun = dchisq, args = list(df = 1), color = "black") +
stat_function(fun = dchisq, args = list(df = 2), color = "orange") +
stat_function(fun = dchisq, args = list(df = 4), color = "cornflowerblue") +
theme_bw()
```

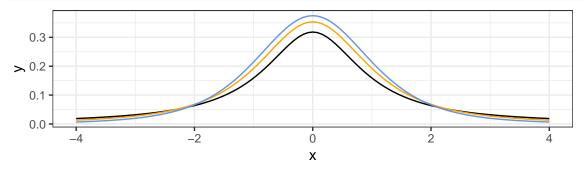


 t_{ν}

A real number.

```
parameters \nu: degrees of freedom  \text{p.f.} \quad f(x|\nu) = \frac{\Gamma\left(\frac{\nu+1}{2}\right)}{\sqrt{\nu\pi}\Gamma\left(\frac{\nu}{2}\right)} \left(1 + \frac{x^2}{\nu}\right)^{\frac{\nu+1}{2}}  Mean 0 \text{ for } \nu > 1, \text{ otherwise undefined} Variance \frac{\nu}{\nu-2} \text{ for } \nu > 2 R functions \text{dt}(\dots, \text{df} = \nu), \text{pt, qt, rt}
```

```
ggplot(data = data.frame(x = c(-4, 4)), mapping = aes(x = x)) +
    stat_function(fun = dt, args = list(df = 1), color = "black") +
    stat_function(fun = dt, args = list(df = 2), color = "orange") +
    stat_function(fun = dt, args = list(df = 4), color = "cornflowerblue") +
    theme_bw()
```



```
\mathbf{F}(\nu_1,\nu_2)
```

A non-negative real number.

```
parameters \nu_1: degrees of freedom, \nu_2: degrees of freedom R functions df(..., df1 = \nu_1, df2 = \nu_2), pf, qf, rf
```

```
ggplot(data = data.frame(x = c(0, 4)), mapping = aes(x = x)) +
    stat_function(fun = df, args = list(df1 = 1, df2 = 1), color = "black") +
    stat_function(fun = df, args = list(df1 = 2, df2 = 7), color = "orange") +
    stat_function(fun = df, args = list(df1 = 4, df2 = 2), color = "cornflowerblue") +
    theme_bw()
```

Χ

$\mathbf{Pareto}(x_0, \alpha)$

A real number that is greater than or equal to x_0 .

```
parameters x_0: lower bound of support, \alpha: shape parameter p.f. f(x|x_0,\alpha) = \frac{\alpha x_0^{\alpha}}{x^{\alpha+1}} \mathbb{I}_{[x_0,\infty)}(x) = \begin{cases} \frac{\alpha x_0^{\alpha}}{x^{\alpha+1}} & \text{if } x \geq x_0 \\ 0 & \text{otherwise} \end{cases} R functions Not provided as part of base R
```

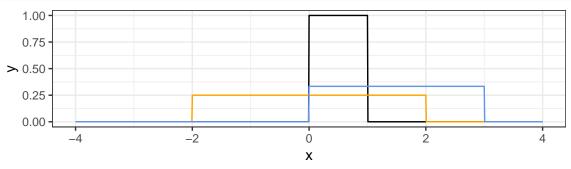
```
#' Calculate pdf of a Pareto distribution
# '
#' Oparam x vector of real numbers at which to evaluate the Pareto density
#' Oparam x_O location parameter for the Pareto distribution
#' Oparam alpha scale parameter for the Pareto distribution
#'
#' Creturn vector of the same length as x of values of the Pareto density function.
dpareto <- function(x, x_0, alpha) {</pre>
    result <- rep(-Inf, length(x))
    inds_x_greater_x_0 <- (x > x_0)
    result[inds_x_greater_x_0] \leftarrow log(alpha) + alpha * log(x_0) - (alpha + 1) * log(x[inds_x_greater_x_0])
    return(exp(result))
}
ggplot(data = data.frame(x = c(0, 20)), mapping = aes(x = x)) +
    stat_function(fun = dpareto, args = c(x_0 = 1, alpha = 1), n = 1001) +
    stat_function(fun = dpareto, args = c(x_0 = 5, alpha = 3), n = 1001, color = "orange") +
    stat_function(fun = dpareto, args = c(x_0 = 5, alpha = 2), n = 1001, color = "cornflowerblue") +
  theme bw()
  1.00
  0.75
> 0.50
  0.25
  0.00
                                            10
                                                             15
                                                                               20
                                            Х
```

Uniform(a, b)

A real number between a and b (inclusive).

```
parameters a: lower endpoint, b: upper endpoint  p.f. \quad f(x|a,b) = \begin{cases} \frac{1}{b-a} & \text{if } x \in [a,b] \\ 0 & \text{otherwise} \end{cases}  Mean  \frac{1}{2}(a+b)  Variance  \frac{1}{12}(b-a)^2  R functions  \frac{1}{2}(a+b) = a, \text{ max } = b, \text{ punif, qunif, runif}
```

```
ggplot(data = data.frame(x = c(-4, 4)), mapping = aes(x = x)) +
stat_function(fun = dunif, args = list(min = 0, max = 1), color = "black", n = 1001) +
stat_function(fun = dunif, args = list(min = -2, max = 2), color = "orange", n = 1001) +
stat_function(fun = dunif, args = list(min = 0, max = 3), color = "cornflowerblue", n = 1001) +
theme_bw()
```



$\mathbf{Beta}(a,b)$

A real number between 0 and 1.

```
parameters a: lower endpoint, b: upper endpoint  p.f. \quad f(x|a,b) = \begin{cases} \frac{\Gamma(a+b)}{\Gamma(a)\Gamma(b)}x^{(a-1)}(1-x)^{(b-1)}, & \text{if } 0 \leq x \leq 1 \\ 0 & \text{otherwise} \end{cases}  Mean \frac{a}{a+b} Variance \frac{ab}{(a+b)^2(a+b+1)} R functions deta(\ldots, min = a, max = b), punif, qunif, runif
```

```
ggplot(data = data.frame(x = c(0, 1)), mapping = aes(x = x)) +
    stat_function(fun = dbeta, args = list(shape1 = 1, shape2 = 1), color = "black") +
    stat_function(fun = dbeta, args = list(shape1 = 0.5, shape2 = 2), color = "orange") +
    stat_function(fun = dbeta, args = list(shape1 = 10, shape2 = 3), color = "cornflowerblue") +
    theme_bw()
```

