

# Package: ggibbous (via r-universe)

September 9, 2024

**Title** Moon Charts, a Pie Chart Alternative

**Version** 0.1.1

**URL** <https://github.com/mnbram/ggibbous>

**BugReports** <https://github.com/mnbram/ggibbous/issues>

**Description** Moon charts are like pie charts except that the proportions are shown as crescent or gibbous portions of a circle, like the lit and unlit portions of the moon. As such, they work best with only one or two groups. 'ggibbous' extends 'ggplot2' to allow for plotting multiple moon charts in a single panel and does not require a square coordinate system.

**Depends** ggplot2 (>= 3.2.1), R (>= 3.6.0)

**Imports** grid, scales (>= 1.0.0)

**Suggests** knitr, mapproj, maps, rmarkdown, testthat (>= 2.1.0), vdiffr (>= 0.3.1)

**License** GPL-3

**LazyData** true

**RoxygenNote** 7.1.1

**Encoding** UTF-8

**VignetteBuilder** knitr

**Repository** <https://mnbram.r-universe.dev>

**RemoteUrl** <https://github.com/mnbram/ggibbous>

**RemoteRef** HEAD

**RemoteSha** 5aed4b08cd5e78aa4dde3f5597eb50580b1ebc1d

## Contents

dmeladh . . . . .	2
draw_key_moon . . . . .	2
geom_moon . . . . .	3
ggibbous . . . . .	5
lunardist . . . . .	6
moonGrob . . . . .	7

**Index****9**


---

dmeladh	<i>Adh allele frequencies in Australasian Drosophila melanogaster</i>
---------	---

---

**Description**

This data set contains allele frequencies for the "fast" and "slow" variants of the enzyme alcohol dehydrogenase in Australasian (mostly Australian) populations of *Drosophila melanogaster*. The data are taken from Oakeshott, J.G., et al. 1982. Alcohol dehydrogenase and glycerol-3-phosphate dehydrogenase clines in *Drosophila melanogaster* on different continents. *Evolution*, 36(1): 86-96.

**Usage**

dmeladh

**Format**

A data frame with 34 rows and 6 variables:

**Locality** location of population sample

**Latitude** latitude of population sample

**Longitude** longitude of population sample

**N** number of samples

**AdhF** percent of samples with *Adh<sup>F</sup>* allele, as an integer

**AdhS** percent of samples with *Adh<sup>S</sup>* allele, as an integer

---

draw_key_moon	<i>Moon key glyph for legends</i>
---------------	-----------------------------------

---

**Description**

Draws the legend key glyphs used in geom\_moon.

**Usage**

draw\_key\_moon(data, params, size)

draw\_key\_moon\_left(data, params, size)

draw\_key\_full\_moon(data, params, size)

**Arguments**

data	A single row data frame containing the scaled aesthetics to display in this key
params	A list of additional parameters supplied to the geom.
size	Width and height of key in mm.

**Details**

draw\_key\_moon (the default in geom\_moon) draws a gibbous moon filled from the right. draw\_key\_moon\_left draws a crescent moon from the right. draw\_key\_full\_moon draws a circle, which is very similar to draw\_key\_point in ggplot2, but the size is calculated slightly differently and the default aesthetics differ.

**Value**

A grid grob.

---

geom\_moon

*Moon charts*

---

**Description**

The moon geom is used to create moon charts, which are like pie charts except that the proportions are shown as crescent or gibbous portions of a circle, like the lit and unlit portions of the moon. As such, they work best with only one or two groups.

**Usage**

```
geom_moon(
  mapping = NULL,
  data = NULL,
  stat = "identity",
  position = "identity",
  na.rm = FALSE,
  show.legend = NA,
  inherit.aes = TRUE,
  ...
)
```

**Arguments**

mapping	Set of aesthetic mappings created by <code>aes()</code> or <code>aes_()</code> . If specified and <code>inherit.aes = TRUE</code> (the default), it is combined with the default mapping at the top level of the plot. You must supply mapping if there is no plot mapping.
---------	---

<code>data</code>	<p>The data to be displayed in this layer. There are three options:</p> <p>If <code>NULL</code>, the default, the data is inherited from the plot data as specified in the call to <code>ggplot()</code>.</p> <p>A <code>data.frame</code>, or other object, will override the plot data. All objects will be fortified to produce a data frame. See <code>fortify()</code> for which variables will be created.</p> <p>A function will be called with a single argument, the plot data. The return value must be a <code>data.frame</code>, and will be used as the layer data. A function can be created from a formula (e.g. <code>~ head(.x, 10)</code>).</p>
<code>stat</code>	The statistical transformation to use on the data for this layer, as a string.
<code>position</code>	Position adjustment, either as a string, or the result of a call to a position adjustment function.
<code>na.rm</code>	If <code>FALSE</code> , the default, missing values are removed with a warning. If <code>TRUE</code> , missing values are silently removed.
<code>show.legend</code>	logical. Should this layer be included in the legends? <code>NA</code> , the default, includes if any aesthetics are mapped. <code>FALSE</code> never includes, and <code>TRUE</code> always includes. It can also be a named logical vector to finely select the aesthetics to display.
<code>inherit.aes</code>	If <code>FALSE</code> , overrides the default aesthetics, rather than combining with them. This is most useful for helper functions that define both data and aesthetics and shouldn't inherit behaviour from the default plot specification, e.g. <code>borders()</code> .
<code>...</code>	Other arguments passed on to <code>layer()</code> . These are often aesthetics, used to set an aesthetic to a fixed value, like <code>colour = "red"</code> or <code>size = 3</code> . They may also be parameters to the paired geom/stat.

## Details

`geom_moon` acts like `geom_point` in that multiple moons can be plotted on the same panel with `x` and `y` in the plot's coordinate system, but size determined independently of the coordinate system. This behavior also means that the moons will always be circular even if the coordinate system is not square.

In order to get a full circle with two complementary sections (a crescent and a gibbous moon), you need to plot two shapes: one with `right = TRUE` and one with `right = FALSE`, with `ratio` on the second one equal to `1 - ratio` on the first.

## Aesthetics

`x` and `y` are required aesthetics. `size`, `fill`, `colo(u)r`, `alpha`, `stroke`, and `group` aesthetics are understood as in other geoms. Two new aesthetics are also introduced: `ratio` and `right`. `ratio` controls the proportion of the moon to be plotted, from 0 to 1. `right` takes a boolean value to indicate whether the moon should be filled from the right or the left.

## Examples

```
ggplot(
  data.frame(x = 1:5, y = 1, size = 1:5, ratio = 1:5 * 0.2),
  aes(x = x, y = y, size = size, ratio = ratio)
) +
```

```

geom_moon()

# To make full moon charts, you need two calls to geom_moon(), one with
# right = TRUE and one with right = FALSE and ratio equal to 1 - ratio
# from the first one
ggplot(dmeladh) +
  geom_moon(
    x = 0.5, y = 0.5, fill = "forestgreen", color = "forestgreen",
    aes(ratio = AdhF / 100)
  ) +
  geom_moon(
    x = 0.5, y = 0.5, fill = "gold", color = "gold",
    aes(ratio = AdhS / 100), right = FALSE
  ) +
  facet_wrap(~Locality, ncol = 7)

# The same thing can be accomplished with a single call to geom_moon()
# using a "long" data frame with both frequencies if you set a grouping
# variable and set the `right` variable to a boolean column
dmeladh_long <- reshape(
  dmeladh,
  varying = c("AdhF", "AdhS"),
  v.names = "freq",
  timevar = "allele",
  times = c("AdhF", "AdhS"),
  idvar = c("Locality", "Latitude", "Longitude", "N"),
  direction = "long"
)
dmeladh_long$right <- rep(c(TRUE, FALSE), each = nrow(dmeladh))
ggplot(dmeladh_long) +
  geom_moon(
    x = 0.5, y = 0.5, key_glyph = draw_key_rect,
    aes(ratio = freq / 100, fill = allele, color = allele, right = right),
  ) +
  facet_wrap(~Locality, ncol = 7)

# Moon charts (and pie charts) are sometimes useful on maps when x and y
# cannot be used as aesthetic dimensions because they are already spatial
# dimensions. Overplotting needs to be considered carefully, however.
ggplot(
  subset(dmeladh, N > 200),
  aes(Longitude, Latitude)
) +
  geom_moon(aes(ratio = AdhF / 100), fill = "black") +
  geom_moon(aes(ratio = AdhS / 100), right = FALSE) +
  coord_fixed()

```

## Description

Moon charts are like pie charts except that the proportions are shown as crescent or gibbous portions of a circle, like the lit and unlit portions of the moon. As such, they work best with only one or two groups. `gggibbous` extends `ggplot2` to allow for plotting multiple moon charts in a single panel and does not require a square coordinate system.

## Details

The workhorse function is `geom_moon`, which adds a moon chart layer to a `ggplot2` plot. The `draw_key_moon`, `draw_key_moon_left`, and `draw_key_full_moon` functions provides legend key glyphs for plots that use `geom_moon`. There are also functions for the raw grid grobs: `grid.moon` and `moonGrob`.

For more information, see the `gggibbous` vignette.

---

lunardist

*Lunar distances and principal phases for 2019*

---

## Description

This data set contains the distance from the Earth to the Moon for each day in 2019, as well as the dates (in UTC) of each occurrence of the four principal phases of the moon. The data are adapted from NASA.

## Usage

```
lunardist
```

## Format

A data frame with 365 rows and 3 variables:

**date** Date

**distance** Distance from the Earth to the Moon in kilometers

**phase** Principal phase of the moon (full, new, first quarter, third quarter) or NA if no principal phase on that date

## Source

<https://svs.gsfc.nasa.gov/4442>

---

moonGrob	<i>Draw a moon</i>
----------	--------------------

---

## Description

Functions to create and draw crescent or gibbous moon shapes.

## Usage

```
moonGrob(
  x,
  y,
  ratio = 0.25,
  right = TRUE,
  r = 10,
  angle = 0,
  default.units = "npc",
  size.units = "mm",
  ...
)

grid.moon(..., draw = TRUE)
```

## Arguments

x	A numeric vector or unit object specifying x-locations.
y	A numeric vector or unit object specifying y-locations.
ratio	A numeric vector with values from 0 to 1 specifying the proportion of the moons to draw.
right	A boolean vector specifying whether the moon should be filled from the right (TRUE) or left (FALSE).
r	A numeric vector specifying the radii of the circles describing the moons.
angle	Not used.
default.units	A string indicating the default units to use if x, y, width, or height are only given as numeric vectors.
size.units	A string indicating the units to use for the radii of the moons.
...	Arguments passed on to <code>grid::polygonGrob</code> .
draw	A logical value indicating whether graphics output should be produced.

## Details

Both functions create a moon grob (a graphical object describing a crescent or gibbous moon), but only `grid.moon` draws the moon (and then only if `draw` is TRUE).

These functions calculate a set of points describing the perimeters of the moons and pass these points on to `grid::polygonGrob`.

The units in `default.units` and `size.units` can be different; `grid` will add them together appropriately before drawing.

**Value**

A grob object.

**Examples**

```
grid::grid.newpage()  
grid.moon(x = 1:3 * 0.25, y = rep(0.5, 3), ratio = 1:3 * 0.25, r = 10)
```



# Index

## \* datasets

dmeladh, [2](#)

lunardist, [6](#)

aes(), [3](#)

aes\_(), [3](#)

borders(), [4](#)

dmeladh, [2](#)

draw\_key\_full\_moon (draw\_key\_moon), [2](#)

draw\_key\_moon, [2](#)

draw\_key\_moon\_left (draw\_key\_moon), [2](#)

fortify(), [4](#)

geom\_moon, [3](#)

gggibbous, [5](#)

ggplot(), [4](#)

grid.moon (moonGrob), [7](#)

layer(), [4](#)

lunardist, [6](#)

moonGrob, [7](#)