GGally Notes

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Contents

ggcorr to plot correlation matrices

1

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Correlation matrices show correlation coefficients between many continuous variables. These matrices are created using the cor function. The ggcorr function from the ggally package to plot correlation matrices.

Usage

```
ggcorr(data, method = "pairwise", palette = "RdYlGn", ...)
data data matrix
method character string giving a method for computing covariances. Defaults to "pairwise".
palette ColorBrewer palette to be used for correlation coefficients. Defaults to "RdYlGn".
geom geom object to use. Accepts either tile (the default) or circle, to plot proportionally scaled circles.
```

```
#install.packages("GGally")
library(GGally)
library(ggplot2)
library(dplyr)
library(tidyr)
library(rdryad)

# retrieve Analis trait data from Kolbe et al. 2011, Evolution 65(12): 3608-3624.
# http://datadryad.org/handle/10255/dryad.34389?show=full
anolis.data <- download_url("10255/dryad.34389")
anolis.traits <- dryad_getfile(anolis.data)
#is.data.frame(anolis.traits)
anolis.traits <- tbl_df(anolis.traits)
glimpse(anolis.traits)</pre>
```

```
## Observations: 21
## Variables: 17
## $ Species
                                     (fctr) A. allisoni, A. alutaceus, A. ...
## $ Ecomorph
                                     (fctr) TC, GB, TW, CG, TC, CG, TG, T,...
## $ Snout.vent.length
                                     (dbl) 76.1300, 35.5000, 42.1250, 134....
                                     (dbl) 24.74000, 10.13750, 12.71250, 3...
## $ Head.length
## $ Head.width
                                     (dbl) 13.63000, 5.15000, 6.31250, 23....
## $ Head.height
                                     (dbl) 9.17500, 4.13750, 4.97500, 20.5...
## $ Pectoral.width
                                     (dbl) 10.05000, 4.20000, 4.90000, 16....
## $ Pelvis.width
                                     (dbl) 7.780000, 2.940000, 4.067500, 1...
## $ Lamellae.3rd.foretoe
                                     (dbl) 21.90000, 11.60000, 11.80000, 2...
                                     (dbl) 30.00000, 17.16700, 17.25000, 3...
## $ Lamellae.4th.hindtoe
```

```
## $ Humerus.length (dbl) 12.545000, 5.195000, 5.980000, ...

## $ Ulna.length (dbl) 8.80925, 3.92000, 3.99000, 18.0...

## $ Phalanx.I.3rd.foretoe.length (dbl) 2.859750, 1.105000, 1.500000, 5...

## $ Fumer.length (dbl) 14.096750, 9.617500, 7.657500, ...

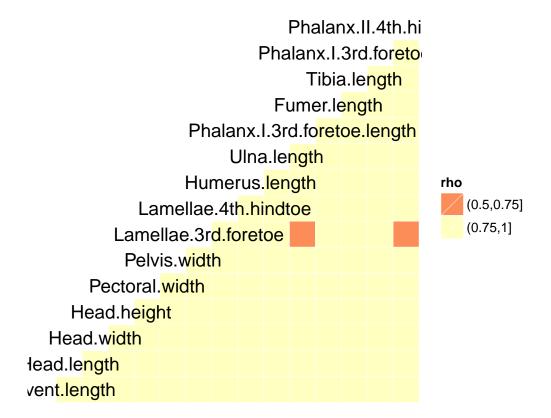
## $ Tibia.length (dbl) 12.565000, 7.980000, 5.892500, ...

## $ Phalanx.I.3rd.foretoe.length.1 (dbl) 7.175750, 5.275000, 3.824250, 1...

## $ Phalanx.II.4th.hindtoe.length (dbl) 5.090000, 2.588000, 2.054500, 8...

select(anolis.traits, -(Species:Ecomorph)) %>%

ggcorr()
```



ggcorr(anolis.traits[, -c(1, 2)])

Phalanx.II.4th.hi Phalanx.I.3rd.foreto Tibia.length Fumer.length Phalanx.I.3rd.foretoe.length Ulna.length Humerus.length rho (0.5,0.75] Lamellae.4th.hindtoe (0.75,1]Lamellae.3rd.foretoe Pelvis.width Pectoral.width Head.height Head.width lead.length ven<mark>t.length</mark>