# 1 ggtreeExtra: A universal package to visualize compact circular layers of phylogenetic tree

Shuangbin Xu, Zehan Dai, Pingfan Guo, Xiaocong Fu, Shanshan Liu, Lang Zhou, Wenli Tang, Tingze Feng, Meijun Chen, Li Zhan and Guangchuang Yu\*

## 1.1 Examples to show how to use *ggtreeExtra* with some geometric function defined by ggplot2-based packages

The supplementary file1 has listed several examples to show how to map the associated data to circular phylogenetic tree using ggtreeExtra. These examples has contained the illustration how to use ggtreeExtra with common geometric function defined by ggplot2(Wickham 2016). Here, we show several cases that ggtreeExtra works with geometric function defined by ggplot2-based packages.

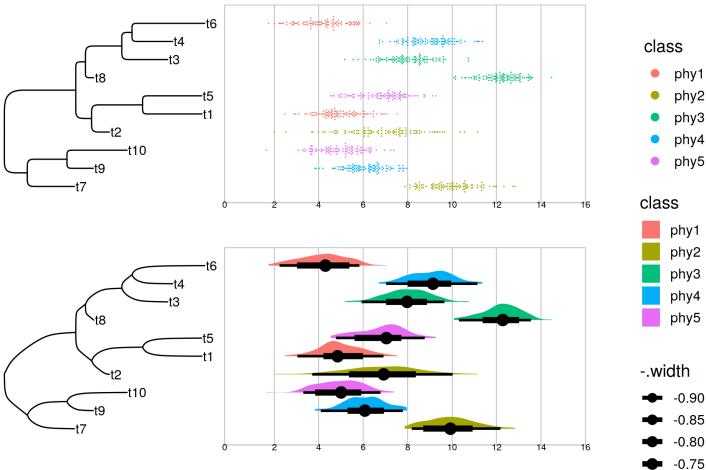
#### 1.1.1 Working with ggdist package

ggdist(Kay 2020) can be used to display the distribution and uncertainty of data, ggtreeExtra can work with several geometric function defined by it (supplementary file Table.S1). If the associated data should be displayed the distribution, such as the gene expression in different samples.

```
library(tibble)
library(tidyr)
library(ggdist)
library(ggtree)
library(ggplot2)
library(ggtreeExtra)
library(patchwork)
set.seed(1024)
df = tribble(
    ~id, ~class, ~value,
    "t1", "phy1", rnorm(100, mean = 5),
    "t2", "phy2", rnorm(100, mean = 7, sd = 1.5),
    "t3", "phy3", rnorm(100, mean = 8),
    "t4",
          "phy4", rnorm(100, mean = 9),
    "t5", "phy5", rnorm(100, mean = 7),
    "t6", "phy1", rnorm(100, mean = 4),
          "phy2", rnorm(100, mean = 10),
    "t7",
          "phy3", rnorm(100, mean = 12),
    "t9", "phy4", rnorm(100, mean = 6),
    "t10", "phy5", rnorm(100, mean = 5)
  ) %>% unnest(value)
tr <- rtree(10)
p1 <- ggtree(tr, layout="roundrect", size=0.3) + geom_tiplab(size=3)
p2 <- ggtree(tr, layout="ellipse", size=0.3) + geom_tiplab(size=3)
# The associate data (df)
p1 <- p1 +
      geom_fruit(data=df,
                 geom=geom dots,
                 mapping=aes(y=id, x=value, fill=class),
                 position=position_identityx(),
                 color=NA,
                 dotsize=3.
```

<sup>\*</sup>correspondence: guangchuangyu@gmail.com, gcyu1@smu.edu.cn

```
pwidth=1.8,
                  offset=0.1,
                  orientation="y",
                  side="both",
                  grid.params=list(),
                  axis.params=list(axis="x",text.size=2, nbreak=6)
      ) + theme(legend.key.size=unit(0.5, "cm"))
p2 <- p2 +
      geom_fruit(data=df,
                  geom=geom_slabinterval,
                  mapping=aes(y=id, x=value, fill=class),
                  position=position_identityx(),
                  pwidth=1.8,
                  offset=0.1,
                  orientation="y",
                  side="right",
                  stat=StatSampleSlabinterval,
                  grid.params=list(),
                  axis.params=list(axis="x",text.size=2, nbreak=6)
      ) +
      theme(legend.key.size=unit(0.5, "cm"))
p1 / p2
                               ·t6
                         -t4
                                                         · · · · › } $>> 88 $9
                                                                                                 class
```



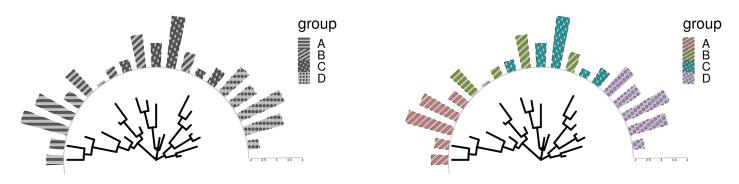
SFig. S1: This example to show ggtreeExtra works with  $geom\_dots$  and  $geom\_slabinterval$  of ggdist(Kay 2020). And ggtreeExtra also support other layout of ggtree (Yu et al. 2017).

#### 1.1.2 Working with ggpattern package

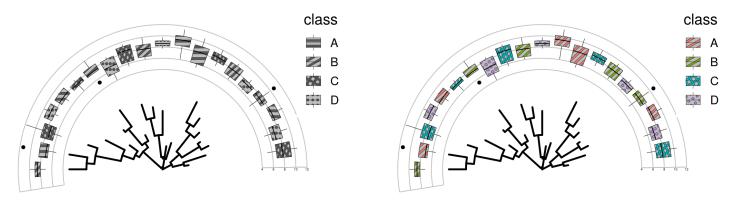
ggpattern(FC~2020) provides geometric layers which support filled areas with geometric and image-based patterns. Some geometric layers of it can also work with ggtreeExtra.

```
library(ggtree)
library(ggtreeExtra)
library(ggpattern)
library(ggplot2)
set.seed(1024)
tr <- rtree(20)
dat <- data.frame(id=tr$tip.label,</pre>
                  value=abs(rnorm(20, 3)),
                  group=c(rep("A", 5), rep("B", 5), rep("C", 5), rep("D", 5)))
dt <- data.frame(id=rep(tr$tip.label, 8), value=abs(rnorm(20 * 8, 8, 1.5)),
                 class=rep(rep(c("A", "B", "C", "D"), 5), 8))
p <- ggtree(
          layout="fan",
          open.angle=180
     )
p1 <- p +
      geom_fruit(
          data=dat,
          geom=geom_bar_pattern,
          mapping=aes(y=id, x=value, pattern=group, pattern_angle=group),
          width=0.6,
          stat="identity",
          pwidth = 0.6,
          pattern_spacing =0.01,
          pattern_size = 0.1,
          pattern_density = 0.4,
          fill = "grey",
          pattern_fill="grey35",
          position=position_identityx(),
          axis.params=list(axis="xy")
      ) +
      theme(
          legend.position= c(0.94, 0.8),
          legend.key.size = unit(0.3, 'cm')
      )
p2 <- p +
      geom_fruit(
          data=dat,
          geom=geom_bar_pattern,
          mapping=aes(y=id, x=value, pattern=group, pattern_fill=group),
          width=0.6,
          stat="identity",
          pwidth = 0.6,
          pattern_spacing =0.01,
          pattern_size = 0.1,
          pattern_density = 0.4,
          fill = "grey",
```

```
position=position_identityx(),
          axis.params=list(axis="xy")
      ) +
      theme(
          legend.position= c(0.94, 0.8),
          legend.key.size = unit(0.3, 'cm')
      )
p3 <- p +
      geom_fruit(
          data=dt,
          geom=geom_boxplot_pattern,
          mapping=aes(y=id, x=value, pattern=class, pattern_angle = class),
          size=0.1,
          outlier.size=0.5,
          pwidth=0.5,
          pattern_size = 0.1,
          pattern_density = 0.4,
          pattern_spacing =0.01,
          fill = "grey",
          pattern_fill="grey35",
          position=position_dodgex(),
          grid.params=list(),
          axis.params=list(axis="x")
      ) +
      theme(
          legend.position= c(0.94, 0.8),
          legend.key.size = unit(0.5, 'cm')
      )
p4 <- p +
      geom_fruit(
          data=dt,
          geom=geom_boxplot_pattern,
          mapping=aes(y=id, x=value, pattern=class, pattern_fill = class),
          size=0.1,
          outlier.size=0.5,
          pwidth = 0.5,
          pattern_size = 0.1,
          pattern_density = 0.4,
          pattern_spacing =0.01,
          fill = "grey",
          position=position_dodgex(),
          grid.params=list(),
          axis.params=list(axis="x")
      ) +
      theme(
          legend.position= c(0.94, 0.8),
          legend.key.size = unit(0.5, 'cm')
      )
p1 + p2
p3 + p4
```



SFig. S2: This example to show ggtreeExtra works with geom\_bar\_pattern and geom\_boxplot\_pattern of ggpattern (FC 2020).



SFig. S3: This example to show ggtreeExtra works with geom\_bar\_pattern and geom\_boxplot\_pattern of ggpattern(FC 2020).

### Reference

FC, Mike. 2020. Ggpattern: Geoms with Patterns.

Kay, Matthew. 2020. ggdist: Visualizations of Distributions and Uncertainty. https://doi.org/10.5281/zenodo.3879620.

Wickham, Hadley. 2016. *Ggplot2: Elegant Graphics for Data Analysis*. Springer-Verlag New York. https://ggplot2.tidy verse.org.

Yu, Guangchuang, David Smith, Huachen Zhu, Yi Guan, and Tommy Tsan-Yuk Lam. 2017. "Ggtree: An R Package for Visualization and Annotation of Phylogenetic Trees with Their Covariates and Other Associated Data." *Methods in Ecology and Evolution* 8 (1): 28–36. https://doi.org/10.1111/2041-210X.12628.