require(ggdendro)

Loading required package: ggdendro

Selección de clústers desde objeto dendrogramas

@param hc objeto hcclust @param k número de clústers

@return @export

@examples

```
dendro_data_k <- function(hc, k) {</pre>
  hcdata
             <- ggdendro::dendro_data(hc, type = "rectangle")</pre>
             <- hcdata$segments</pre>
  seg
  labclust <- cutree(hc, k)[hc$order]</pre>
  segclust <- rep(OL, nrow(seg))</pre>
            <- sort(hc$height, decreasing = TRUE)</pre>
  heights
             <- mean(c(heights[k], heights[k - 1L]), na.rm = TRUE)</pre>
  height
  for (i in 1:k) {
               hcdata$labels$x[labclust == i]
    хi
            <- (seg$x
    idx1
                          >= min(xi)) & (seg$x
                                                      \leq \max(xi)
            <- (seg$xend >= min(xi)) & (seg$xend <= max(xi))</pre>
    idx2
    idx3
            <- seg$yend < height</pre>
            <- idx1 & idx2 & idx3
    idx
    segclust[idx] <- i</pre>
  }
  idx
                           <- which(segclust == 0L)</pre>
  segclust[idx]
                           <- segclust[idx + 1L]</pre>
  hcdata$segments$clust <- segclust
  hcdata$segments$line <- as.integer(segclust < 1L)</pre>
  hcdata$labels$clust
                           <- labclust
  hcdata
}
```

Colocar Etiquetas de Dendrograma

```
Función de creación de dendrograma gráfico con clústers
```

```
"'r plot_ggdendro <- function(hcdata, direction = c("lr", "rl", "tb", "bt"), fan = FALSE, scale.color = NULL, branch.size = 1, label.size = 3, nudge.label = 0.01, expand.y = 0.1) { direction <- match.arg(direction) # if fan = FALSE ybreaks <- pretty(segment(hcdata)y, n = 5)ymax < -max(segment(hcdata)y) ## Ramas p <- ggplot() + geom_segment(data = segment(hcdata), aes(x = x, y = y, xend = xend, yend = yend, linetype = factor(line), colour = factor(clust)), lineend = "round", show.legend = FALSE, size = branch.size)
```

[@]param hcdata @param direction @param fan @param scale.color @param branch.size @param label.size @param nudge.label @param expand.y

Colocar Etiquetas de Dendrograma

```
  \#\# \ Orientación \ if \ (fan) \ \{ \ p < -p + coord\_polar(direction = -1) + scale\_x\_continuous(breaks = NULL, limits = c(0, nrow(label(hcdata)))) + scale\_y\_reverse(breaks = ybreaks) \ \} \ else \ \{ \ p < -p + scale\_x\_continuous(breaks = NULL) \ if \ (direction \%in\% \ c("rl", "lr")) \ \{ \ p < -p + coord\_flip() \ \} \ if \ (direction \%in\% \ c("bt", "lr")) \ \{ \ p < -p + scale\_y\_reverse(breaks = ybreaks) \ \} \ else \ \{ \ p < -p + scale\_y\_continuous(breaks = ybreaks) \ \} \ direction \ (nudge.label) \ \} \ \# \ Etiquetas \ labelParams < -set\_labels\_params(nrow(hcdatalabels), direction, fan)hcdatalabelsangle < -labelParamsangle \ p < -p + geom\_text(data = label(hcdata), aes(x = x, y = y, label = label, colour = factor(clust), angle = angle), vjust = labelParamsvjust, hjust = labelParamshjust, nudge\_y = ymax * nudge.label, size = label.size, show.legend = FALSE) \ \# \ colors \ and \ limits \ if \ (!is.null(scale.color)) \ \{ \ p < -p + scale\_color\_manual(values = scale.color) \ \} \ ylim < -round(ymax * expand.y, 1) \ p < -p + expand\_limits(y = ylim) \ p \ \}  ""
```

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