

```

USarrest.R <- function(){
# Agglomerative clustering algorithms for US Arrest Data
# Requires library "factoextra"

#read data
data("USArrests")
# Load libraries
library("factoextra")

# standardize data
df <- scale(USArrests)
# compute matrix of distances
res.dist <- dist(df, method = "euclidean")

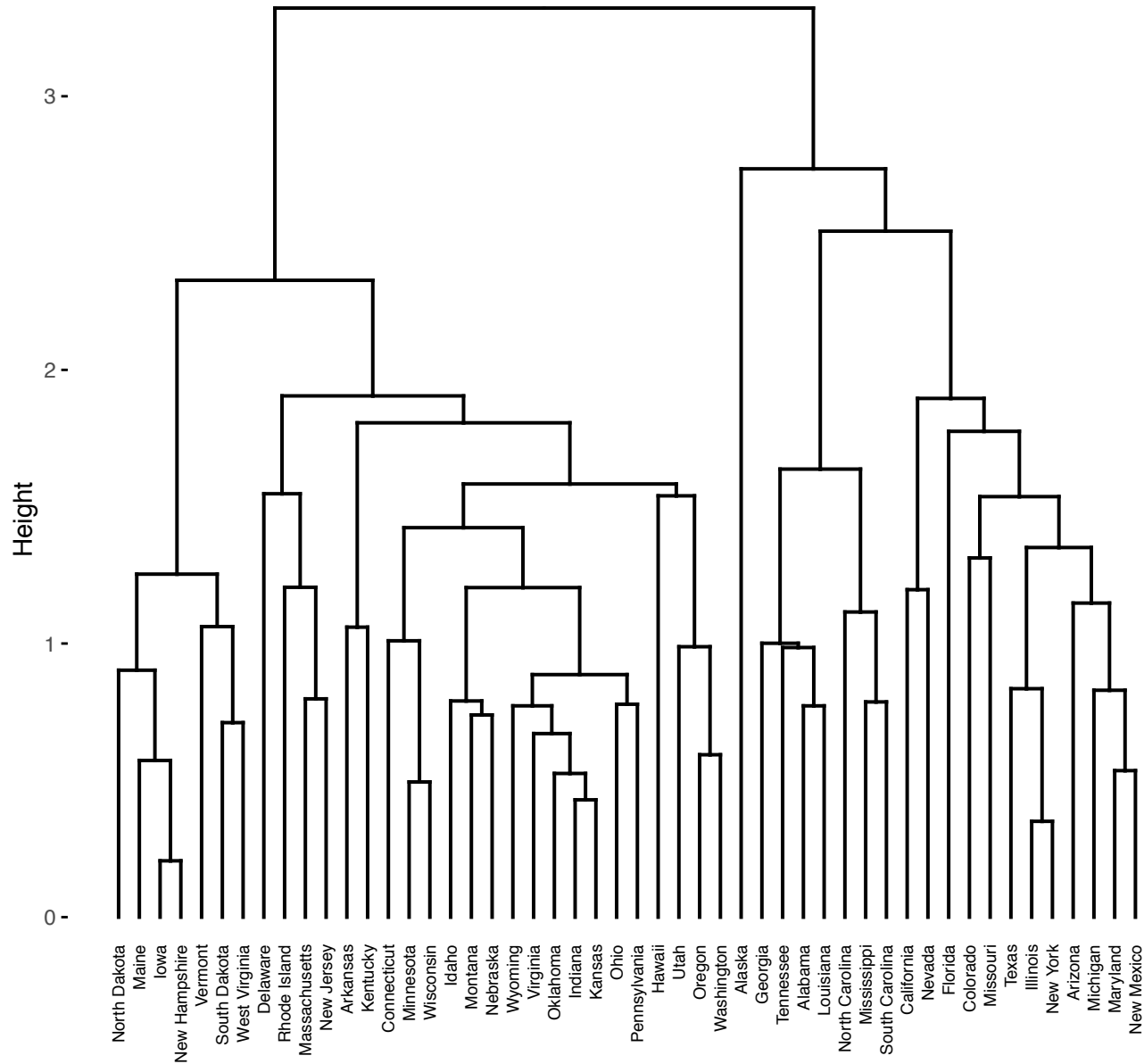
# compute cluster
#res.hc <- hclust(d = res.dist, method = "ward.D2")
#res.hc1 <- hclust(d = res.dist, method = "average")
#res.hc2 <- hclust(d = res.dist, method = "complete")
res.hc3 <- hclust(d = res.dist, method = "single")

# display dendrogram
#plot.hc <- fviz_dend(res.hc, cex = 0.5)
#plot.hc <- fviz_dend(res.hc1, cex = 0.5)
#plot.hc <- fviz_dend(res.hc2, cex = 0.5)
plot.hc <- fviz_dend(res.hc3, cex = 0.5)

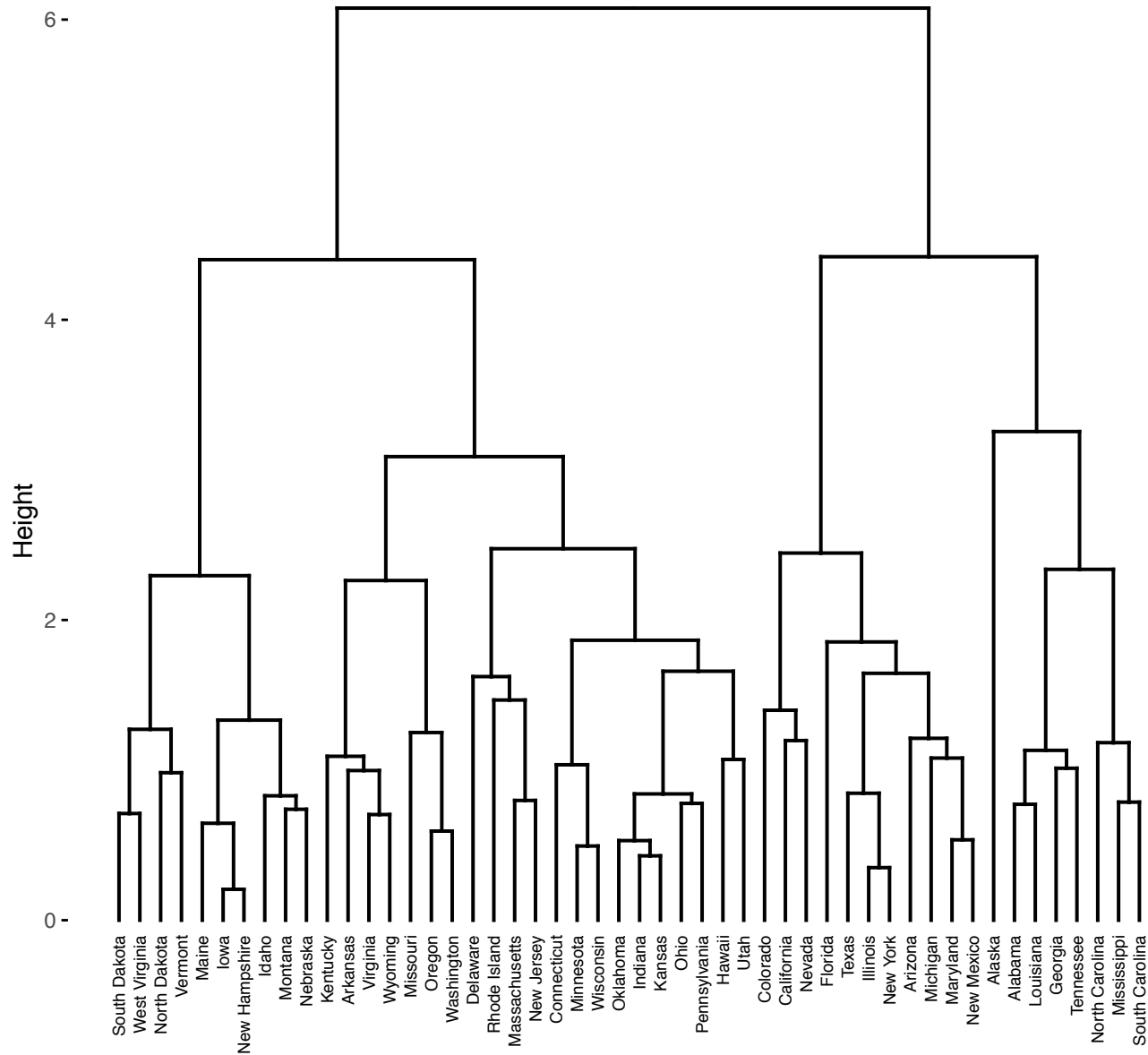
#ggsave("dendogram-ward2.pdf", device = "pdf")
#ggsave("dendogram-average-linkage.pdf", device = "pdf")
#ggsave("dendogram-complete-linkage.pdf", device = "pdf")
ggsave("dendogram-single-linkage.pdf", device = "pdf")
}

```

Cluster Dendrogram



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