

Interactive Phylogenetic Tree

2024-11-19

```
#PACKAGE LOADING
```

```
#Load packages
```

```
library(phytools) #Read in phylogeny
```

```
## Loading required package: ape
```

```
## Loading required package: maps
```

```
library(ape) #For the phylo.vcv function
```

```
library(geiger) #For name.check function
```

```
library(tidyverse) #For piping command
```

```
## -- Attaching core tidyverse packages ----- tidyverse 2.0.0 --
```

```
## v dplyr      1.1.4      v readr      2.1.5
```

```
## v forcats    1.0.0      v stringr    1.5.1
```

```
## v ggplot2    3.5.1      v tibble     3.2.1
```

```
## v lubridate  1.9.3      v tidyr      1.3.1
```

```
## v purrr      1.0.2
```

```
## -- Conflicts ----- tidyverse_conflicts() --
```

```
## x dplyr::filter() masks stats::filter()
```

```
## x dplyr::lag()     masks stats::lag()
```

```
## x purrr::map()     masks maps::map()
```

```
## x dplyr::where()   masks ape::where()
```

```
## i Use the conflicted package (<http://conflicted.r-lib.org/>) to force all conflicts to become errors
```

```
library(ggraph)
```

```
library(igraph)
```

```
##
```

```
## Attaching package: 'igraph'
```

```
##
```

```
## The following objects are masked from 'package:lubridate':
```

```
##
```

```
##      %--%, union
```

```
##
```

```
## The following objects are masked from 'package:dplyr':
```

```
##
```

```
##      as_data_frame, groups, union
```

```
##
```

```
## The following objects are masked from 'package:purrr':
```

```
##
```

```
##      compose, simplify
```

```
##
```

```
## The following object is masked from 'package:tidyr':
```

```
##
```

```
##      crossing
```

```

##
## The following object is masked from 'package:tibble':
##
##   as_data_frame
##
## The following objects are masked from 'package:ape':
##
##   degree, edges, mst, ring
##
## The following objects are masked from 'package:stats':
##
##   decompose, spectrum
##
## The following object is masked from 'package:base':
##
##   union
library(htmlwidgets)
library(phyloanvas)

##
## Attaching package: 'phyloanvas'
##
## The following object is masked from 'package:igraph':
##
##   as_tree

#DATAFRAME
df <- read.csv("~/Desktop/Menstrual taboos/R CODE DATA FILES (USE)/Master's data set.csv", header=TRUE)

#PHYLOGENETIC TREE
phylo<-readNexus("~/Desktop/Menstrual taboos/R CODE DATA FILES (USE)/Time-calibrated SCCS supertree ult.

#MAKE CULTURE NAMES THE ROWNAMES IN DATAFRAME
rownames(df) <- df$tip.label #Set rownames in data frame

pop <- df[,2] #Store the population column

str(df) #Look at data frame

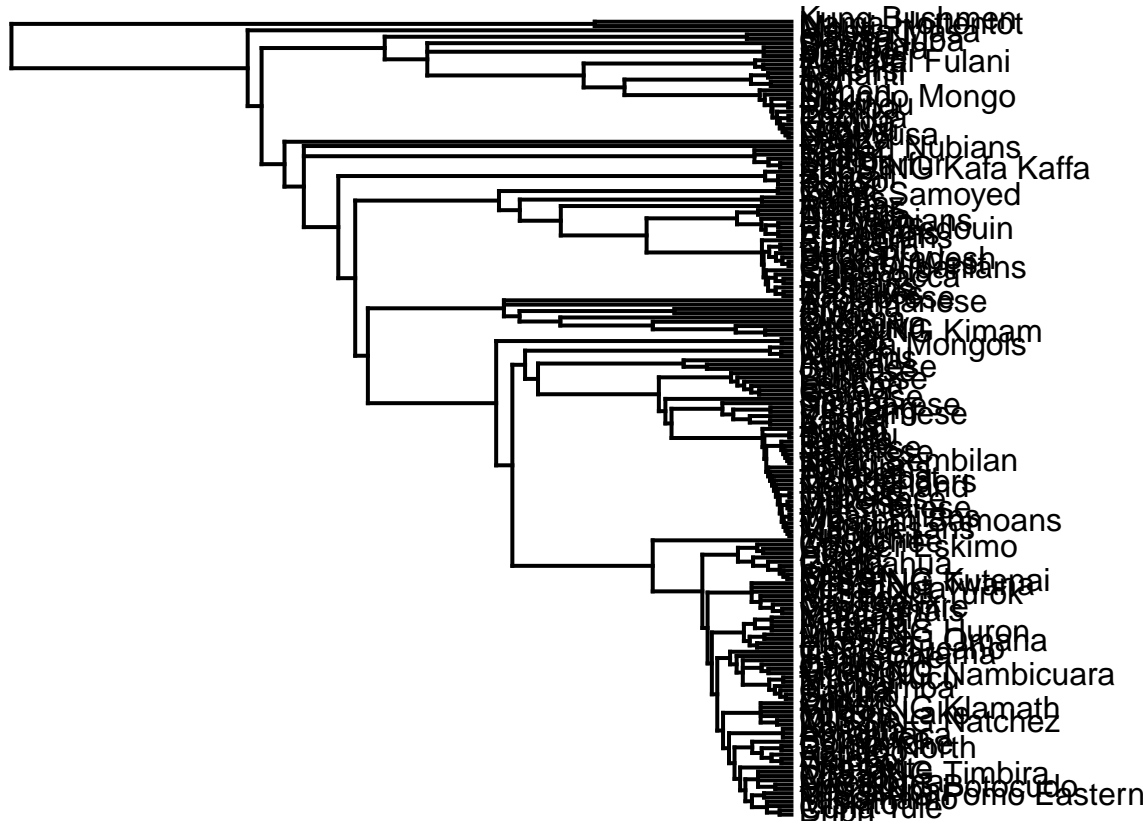
## 'data.frame':   144 obs. of  17 variables:
##  $ HRAF.culture      : chr  "Babylonians" "Zuni" "Teda" "O'odham" ...
##  $ tip.label         : chr  "Babylonians" "Zuni" "Teda" "Papago" ...
##  $ Expression        : int  1 0 1 1 1 1 1 1 1 1 ...
##  $ Restrictiveness   : int  3 0 1 3 1 2 1 3 3 1 ...
##  $ lat               : int  33 36 21 32 23 -24 32 43 23 12 ...
##  $ long              : int  45 -109 18 -112 39 134 -110 -119 7 -72 ...
##  $ Subsistence_Pattern : chr  "Intensive Agricultralists" "Intensive Agricultralists" "Agro-I
##  $ Subsistence       : chr  "Agriculture" "Agriculture" "Agriculture" "Agriculture" ...
##  $ Biome              : chr  "Deserts & Xeric Shrublands" "Deserts & Xeric Shrublands" "Des
##  $ Landscape         : chr  "Deserts" "Deserts" "Deserts" "Deserts" ...
##  $ MeanMonthlyTemperature : int  23 10 22 22 27 21 14 7 20 26 ...
##  $ Temperature_Predictability : num  0.64 0.579 0.642 0.615 0.725 ...
##  $ Temperature_Variance : int  111 103 46 62 24 37 70 84 55 1 ...

```

```
## $ Precipitation_Variance      : num  1.72e+08 7.89e+08 1.57e+08 7.05e+08 1.85e+08 ...
## $ MeanMonthlyPrecipitation    : num   7853 41599 5593 24750 6403 ...
## $ Precipitation_Predictability: num   0.347 0.556 0.304 0.404 0.249 ...
## $ Men_Poly                    : int    0 0 10 27 NA 60 18 12 0 40 ...
```

#CLEAN PHYLOGENETIC TREE FOR ANALYSIS

```
plotTree(phylo) #Look at phylogeny
```



```
name.check (phylo,df$tip.label) # check which populations are in the tree but not in the data and remove
```

```
## $tree_not_data
## [1] "Abipon"           "Abkhaz"           "Ainu"
## [4] "Ajie"             "Aleut"            "Alorese"
## [7] "Amahuaca"         "Amhara"           "Andamanese"
## [10] "Aranda"           "Armenians"        "Ashanti"
## [13] "Atayal"           "Aweikoma"         "Aymara"
## [16] "Azande"           "Aztec"            "Babylonians"
## [19] "Badjau"           "Balinese"         "Bambara"
## [22] "Banen"            "Basques"          "Basseri"
## [25] "Bellacoola"       "Bemba"            "Bogo"
## [28] "Bribri"           "Burmese"          "Buruscho"
## [31] "Callinago"        "Carib_Barama"     "Cayapa"
## [34] "Cayua"            "Chinese"          "Chiricahua"
## [37] "Chukchee"         "Comanche"         "Copper_Eskimo"
## [40] "Creek"            "Cubeo_Tucano"     "Cuna_Tule"
## [43] "Egyptians"        "Eyak"             "Fon"
## [46] "Fur_Darfur"       "Ganda"            "Garos"
## [49] "Gheg_Albanians"   "Gilbertese"       "Gilyak"
```

## [52]	"Goajiro"	"Gond"	"Gros_Ventre"
## [55]	"Hadza"	"Haida"	"Haitians"
## [58]	"Hausa"	"Havasupai"	"Hebrews"
## [61]	"Hidatsa"	"Huichol"	"Iban"
## [64]	"Ibo"	"Ifugao"	"Inca"
## [67]	"Ingalik"	"Irish"	"Japanese"
## [70]	"Javanese"	"Jivaro"	"Kapauku"
## [73]	"Kaska"	"Kazak"	"Kenuzi_Nubians"
## [76]	"Khalka_Mongols"	"Khmer"	"Kikuyu"
## [79]	"Konso"	"Koreans"	"Kung_Bushmen"
## [82]	"Kurd"	"Kwoma"	"Lakher"
## [85]	"Lamet"	"Lapps"	"Lengua"
## [88]	"Lepcha"	"Lolo"	"Lozi"
## [91]	"Luguru"	"Manchu"	"Manus"
## [94]	"Mao"	"Maori"	"Mapuche"
## [97]	"Marquesans"	"Marshallese"	"Masai"
## [100]	"Massa_Masa"	"Mbau_Fijians"	"Mbundu"
## [103]	"Mbuti"	"Mende"	"Micmac"
## [106]	"Miskito"	"MISSING_Botocudo"	"MISSING_Huron"
## [109]	"MISSING_Kafa_Kaffa"	"MISSING_Kimam"	"MISSING_Klamath"
## [112]	"MISSING_Kutenai"	"MISSING_Nambicuara"	"MISSING_Natchez"
## [115]	"MISSING_Omaha"	"MISSING_Pomo_Eastern"	"MISSING_Timbira"
## [118]	"MISSING_Twana"	"MISSING_Yurok"	"Montagnais"
## [121]	"Mundurucu"	"Nama_Hottentot"	"Negri_Sembilan"
## [124]	"New_Ireland"	"Nicobarese"	"Nkundo_Mongo"
## [127]	"Nyakyusa"	"Orokaiva"	"Otoro_Nuba"
## [130]	"Paiute_North"	"Palauans"	"Papago"
## [133]	"Pastoral_Fulani"	"Pawnee"	"Pentecost"
## [136]	"Popoluca"	"Punjabi_West"	"Quiche"
## [139]	"Rshade"	"Riffians"	"Romans"
## [142]	"Russians"	"Rwala_Bedouin"	"Santal"
## [145]	"Saramacca"	"Saulteaux"	"Semang"
## [148]	"Shavante"	"Shilluk"	"Siamese"
## [151]	"Siriono"	"Siuai"	"Slave"
## [154]	"Somali"	"Songhai"	"Suku"
## [157]	"Tallensi"	"Tanala"	"Teda"
## [160]	"Tehuelche"	"Thonga"	"Tikopia"
## [163]	"Tiv"	"Tiwi"	"Tobelorese"
## [166]	"Toda"	"Toradja"	"Trobrianders"
## [169]	"Trukese"	"Trumai"	"Tuareg"
## [172]	"Tupinamba"	"Turks"	"Uttar_Pradesh"
## [175]	"Vedda"	"Vietnamese"	"Warrau"
## [178]	"Western_Samoans"	"Wolof"	"Yahgan"
## [181]	"Yanomamo"	"Yapese"	"Yokuts_Lake"
## [184]	"Yukaghir"	"Yurak_Samoyed"	"Zuni"

```
## $data_not_tree
## NULL
```

```
MT_Tree <- drop.tip (phylo, tip=c("Abipon", "Abkhaz", "Ajie", "Atayal", "Aweikoma", "Banen", "Basseri",
name.check (MT_Tree, df) # OK
```

```
## [1] "OK"
```

