test isOrigin.Oct.9.2018

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Oct. 9, 2018

Install isOrigin package from Github

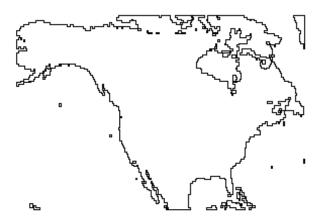
devtools::install_github("SPATIAL-Lab/isorig", force=T)

Load library

```
library(isOrigin)
## Loading required package: raster
## Loading required package: sp
## Loading required package: ggplot2
```

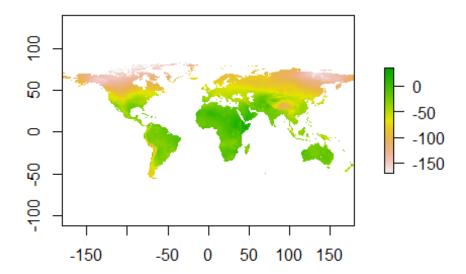
Load North America mask

```
data("naMap")
plot(naMap)
```



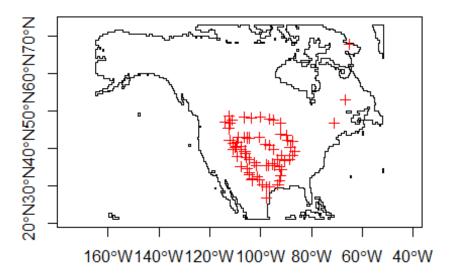
Load world precipitation hydrogen isoscape

data("d2h_world")
plot(d2h_world)



Load hydrogen isotope for human hair in North America

d = subOrigData(taxon = c("Homo sapiens"), mask = naMap)



233 data points are found

```
Exclude some outliers. This step is optional, which depends on your data quality
d <-as.data.frame(d)
dd = d[d$coords.x1<(-80),]
dd <- SpatialPointsDataFrame(dd[,2:3], as.data.frame(dd[,1]))
crs(dd) <- "+proj=longlat +datum=WGS84 +no_defs +ellps=WGS84 +towgs84=0,0,0"</pre>
```

Rescale from environmental isoscape to tissue isoscape

```
r = calRaster(known = dd, isoscape = d2h world, mask = naMap)
##
##
## rescale function uses linear regression model, the summary of this model
is:
##
## Call:
## lm(formula = tissue.iso ~ isoscape.iso)
##
## Residuals:
                1Q Median
      Min
                                3Q
                                       Max
                             5.056 20.184
## -37.317 -4.021
                     0.702
##
```

```
## Coefficients:

## Estimate Std. Error t value Pr(>|t|)

## (Intercept) -71.03824   1.59709   -44.48   <2e-16 ***

## isoscape.iso   0.52349   0.02696   19.42   <2e-16 ***

## ---

## Signif. codes:   0 '***'   0.001 '**'   0.01 '*'   0.05 '.'   0.1 ' ' 1

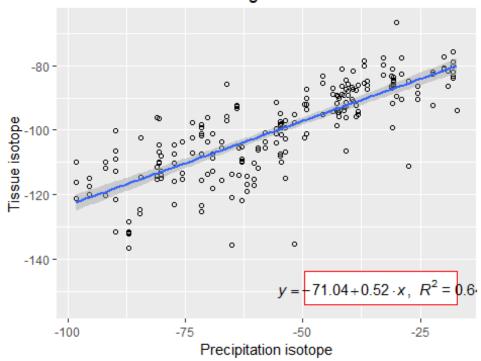
##

## Residual standard error: 8.432 on 213 degrees of freedom

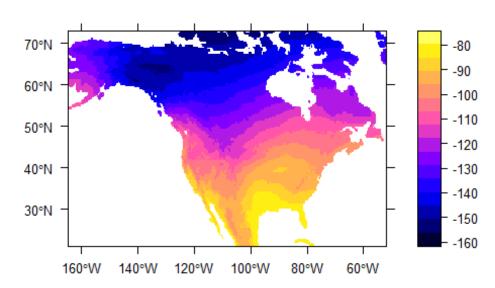
## Multiple R-squared:   0.639, Adjusted R-squared:   0.6373

## F-statistic: 377.1 on 1 and 213 DF, p-value: < 2.2e-16
```

Rescale regression model

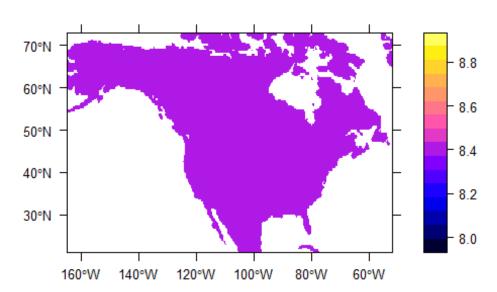


rescale mean



Warning in dir.create("output"): 'output' already exists

rescale sd



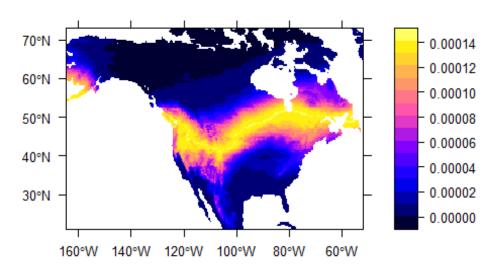
Four unknown-origin examples

```
id = c("A", "B", "C", "D")
d2H = c(-110, -180, -130, -150)
un = data.frame(id,d2H)
```

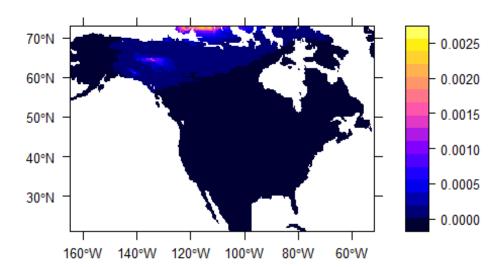
Assignment for unknown-origin examples

```
asn = pdRaster(r$isoscape.rescale,unknown=un,mask=naMap)
## Warning in dir.create("output"): 'output' already exists
## Warning in dir.create("output/pdRaster_Gtif"): 'output\pdRaster_Gtif'
## already exists
```

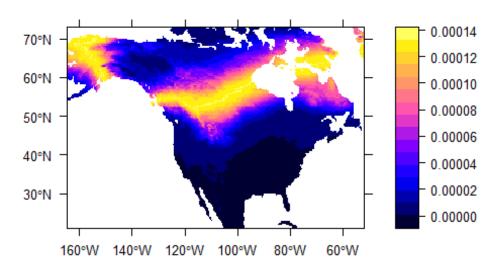
Probability Density Surface for A



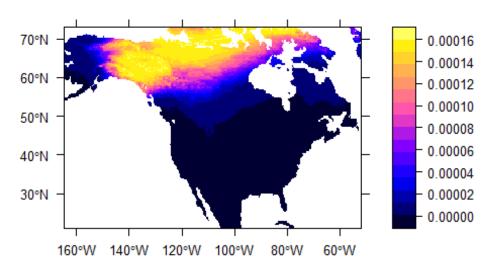
Probability Density Surface for B



Probability Density Surface for C



Probability Density Surface for D



Create SpatialPolygons with two polygons

```
p1 <- c(-100,60,-100,65,-110,65,-110,60,-100,60)
p1 <- matrix(p1, 5,2, byrow = T)
```

p1 <- Polygon(p1)</pre>

```
p1 <- Polygons(list(p1), "p1")
p2 <- c(-100,40,-100,45,-110,45,-110,40,-100,40)
p2 <- matrix(p2, 5,2, byrow = T)
p2 <- Polygon(p2)
p2 <- Polygons(list(p2), "p2")
p12 <- SpatialPolygons(list(p1,p2),1:2)
plot(p12)</pre>
```

Create data.frame with two points

```
pp1 <- c(-100,45)
pp2 <- c(-100,60)
pp12 <- as.data.frame(rbind(pp1,pp2))</pre>
```

Caculate odds ratio for the two polygons created above

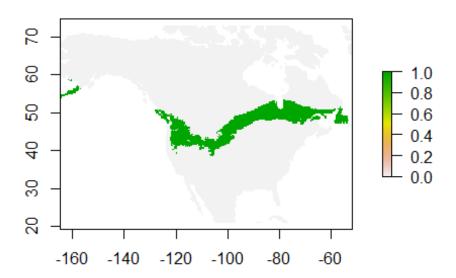
Caculate odds ratio for the two points created above oddsRatio(asn, pp12)

```
## $`P1/P2_odds_ratio`
## A B C D
## 9.078822e+01 2.153343e-10 4.330494e-02 2.066233e-05
##
## $`odds of a pixel to the odds of the max/min pixel`
## ratioToMax.A ratioToMax.B ratioToMax.C ratioToMax.D ratioToMin.A
## 1 0.977370353 1.748220e-13 0.03317593 5.535731e-06 3.638540e+06
## 2 0.000593979 6.746872e-04 0.04225933 2.226383e-01 6.065034e+25
## ratioToMin.B ratioToMin.C ratioToMin.D
## 1 9.199045e-06 1.235069e+05 2.912874e+02
## 2 8.208613e+11 4.315040e+27 2.708740e+14
```

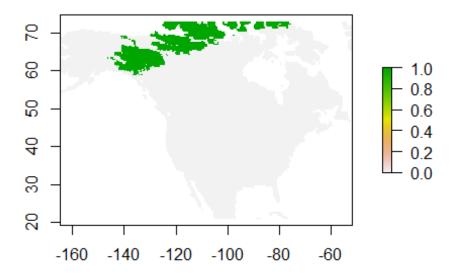
Binary reclassification

```
Top 10% of probability surface (defined by % area)
qtlRaster(asn, threshold = 0.1, thresholdType = 2)
```

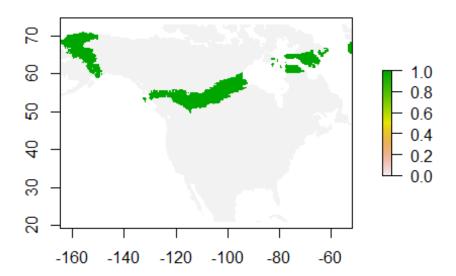
Top 10% by Areafor A



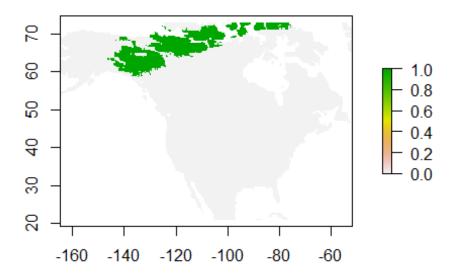
Top 10% by Areafor B



Top 10% by Areafor C



Top 10% by Areafor D



```
## class : RasterStack
## dimensions : 156, 339, 52884, 4 (nrow, ncol, ncell, nlayers)
```

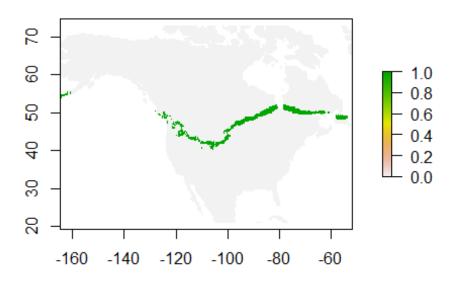
resolution : 0.333, 0.333 (x, y)

extent : -164.682, -51.795, 21.06252, 73.01052 (xmin, xmax, ymin,

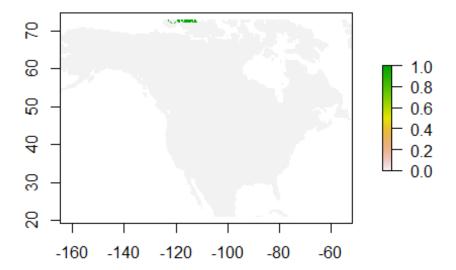
```
ymax)
## coord. ref. : +proj=longlat +datum=WGS84 +no_defs +ellps=WGS84
+towgs84=0,0,0
## names : A, B, C, D
## min values : 0, 0, 0, 0
## max values : 1, 1, 1, 1
```

```
Top 10% of probability surface (defined by % cumulative probability)
qtlRaster(asn, threshold = 0.1, thresholdType = 1)
```

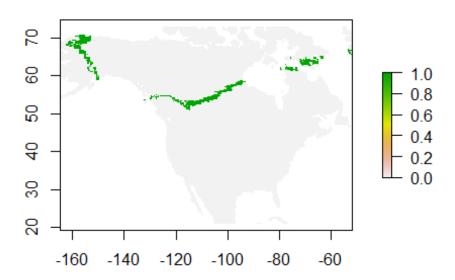
Top 10% by Cumulative Probabilityfor A



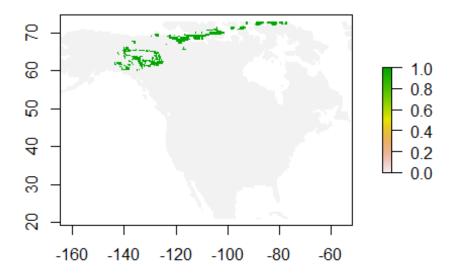
Top 10% by Cumulative Probabilityfor B



Top 10% by Cumulative Probabilityfor C



Top 10% by Cumulative Probabilityfor D

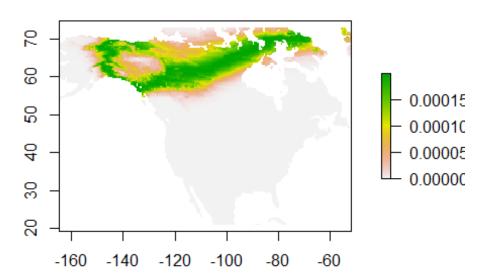


```
## class : RasterStack
## dimensions : 156, 339, 52884, 4 (nrow, ncol, ncell, nlayers)
## resolution : 0.333, 0.333 (x, y)
## extent : -164.682, -51.795, 21.06252, 73.01052 (xmin, xmax, ymin,
```

```
ymax)
## coord. ref. : +proj=longlat +datum=WGS84 +no_defs +ellps=WGS84
+towgs84=0,0,0
## names : A, B, C, D
## min values : 0, 0, 0, 0
## max values : 1, 1, 1, 1
```

Joint probability for individuals of common origin jointP(asn)

Joint Probability

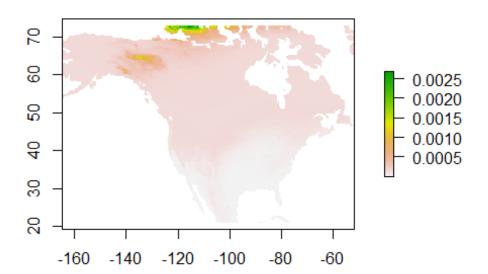


```
## class : RasterLayer
## dimensions : 156, 339, 52884 (nrow, ncol, ncell)
## resolution : 0.333, 0.333 (x, y)
## extent : -164.682, -51.795, 21.06252, 73.01052 (xmin, xmax, ymin, ymax)
## coord. ref. : +proj=longlat +datum=WGS84 +no_defs +ellps=WGS84
+towgs84=0,0,0
## data source : in memory
## names : Joint_Probability
## values : 1.416425e-52, 0.0001997866 (min, max)
```

Probability that at least one individual came from the location (union of probabilities)

unionP(asn)

Union Probability



```
## class : RasterLayer
## dimensions : 156, 339, 52884 (nrow, ncol, ncell)
## resolution : 0.333, 0.333 (x, y)
## extent : -164.682, -51.795, 21.06252, 73.01052 (xmin, xmax, ymin, ymax)
## coord. ref. : +proj=longlat +datum=WGS84 +no_defs +ellps=WGS84
+towgs84=0,0,0
## data source : in memory
## names : layer
## values : 2.239529e-07, 0.002684159 (min, max)
```