# test assignR.20190812

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# Install assignR package from Github

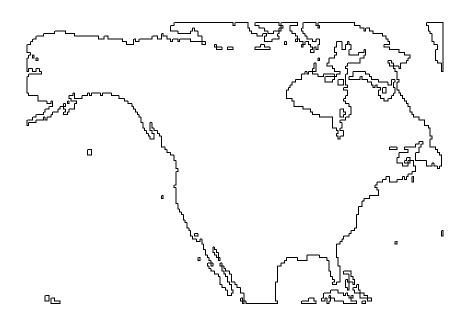
devtools::install\_github("SPATIAL-Lab/assignR", force=T)

# **Load library**

library(assignR)

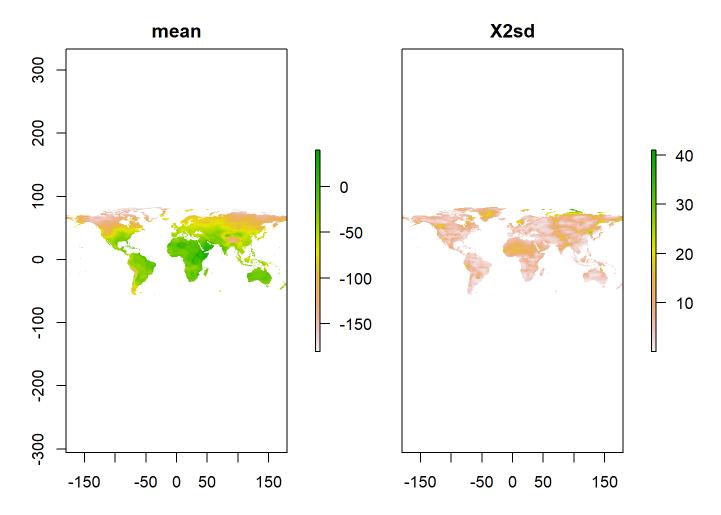
#### **Load North America mask**

data("naMap")
plot(naMap)



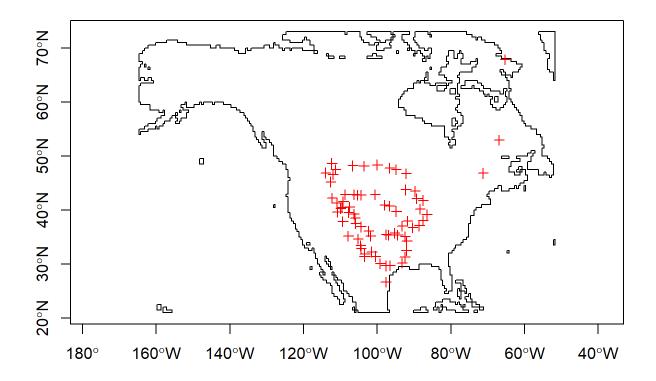
# Load world precipitation hydrogen isoscape

data("d2h\_world")
plot(d2h\_world)



# Load hydrogen isotope for human hair in North America

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

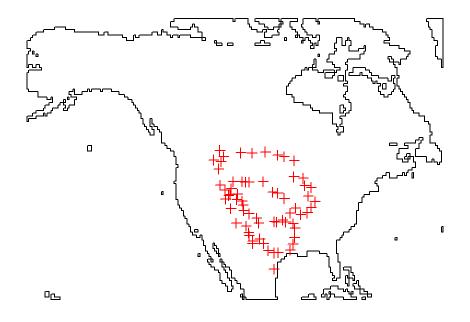


## 233 data points are found

# 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"

plot(naMap)
plot(dd, add=T, col=2)</pre>
```

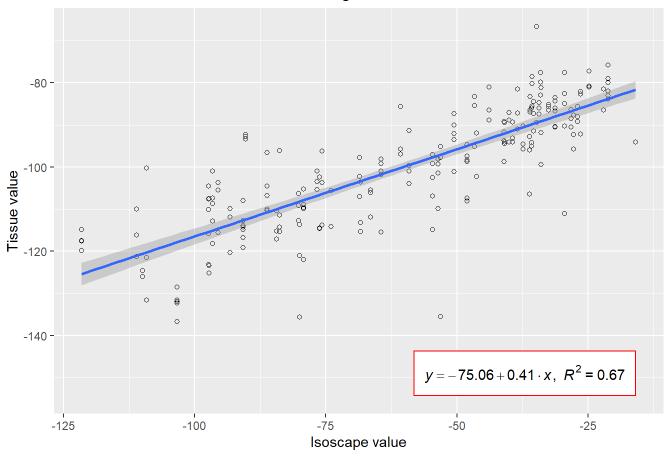


# Rescale from environmental isoscape to tissue isoscape

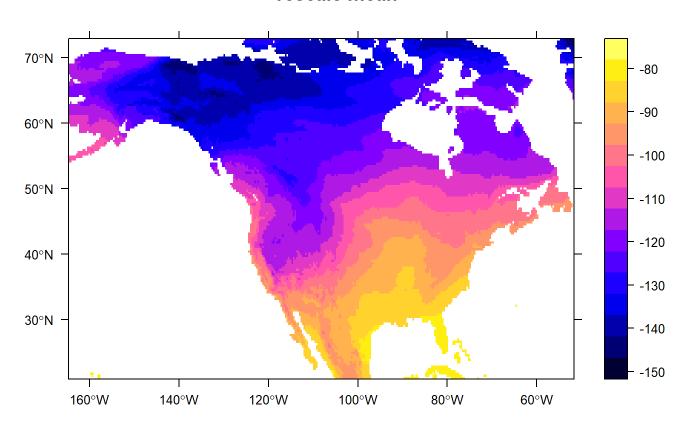
```
r = calRaster(known = dd, isoscape = d2h world, mask = naMap)
## rescale function uses linear regression model, the summary of this
model is:
##
## lm(formula = tissue.iso ~ isoscape.iso[, 1])
## Residuals:
##
   Min
             1Q Median
                           3Q
## -38.407 -4.035 0.233 4.419 22.834
## Coefficients:
##
                   Estimate Std. Error t value Pr(>|t|)
## (Intercept) -75.06067 1.32160 -56.80 <2e-16 ***
                              0.01996 20.74 <2e-16 ***
## isoscape.iso[, 1] 0.41389
```

```
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 8.078 on 213 degrees of freedom
## Multiple R-squared: 0.6687, Adjusted R-squared: 0.6672
## F-statistic: 430 on 1 and 213 DF, p-value: < 2.2e-16</pre>
```

### Rescale regression model

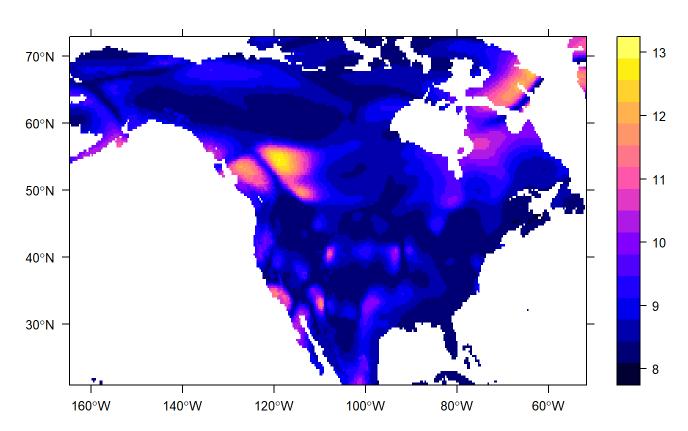


#### rescale mean



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

#### rescale sd



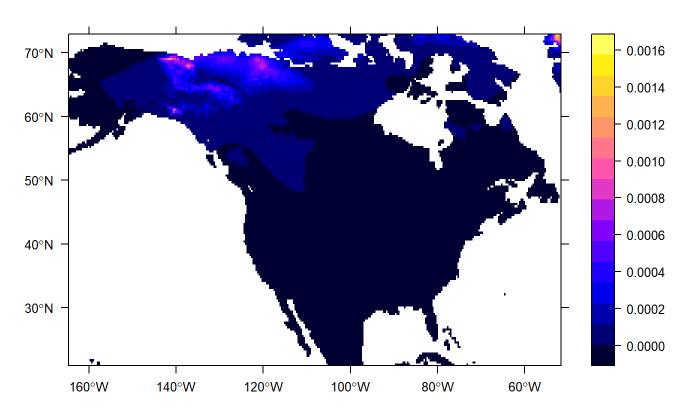
# Four unknown-origin examples

```
id = letters[1:6]
d2H = seq(-160, -80, by=80/5)
un = data.frame(id,d2H)
```

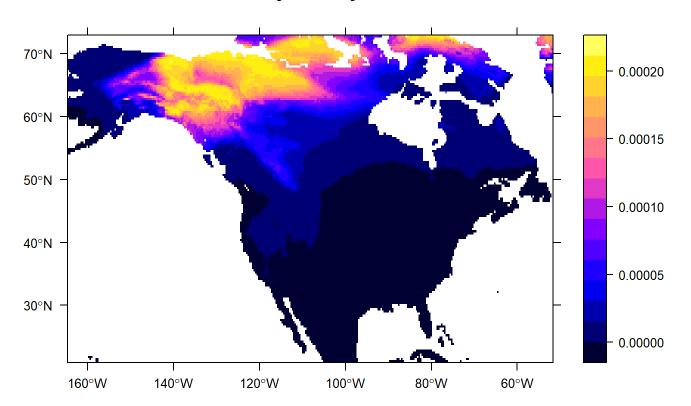
# Assignment for unknown-origin examples

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

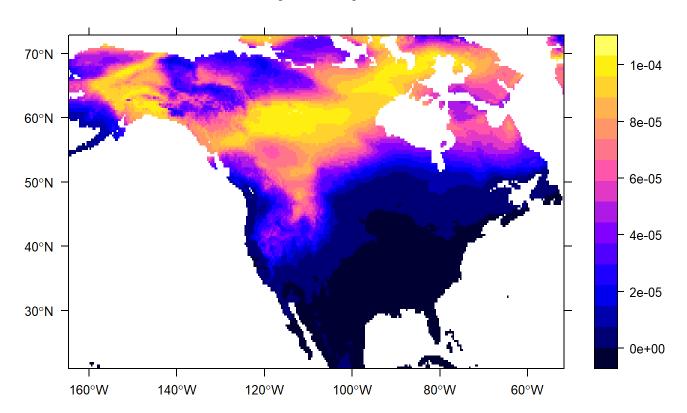
# **Probability Density Surface for a**



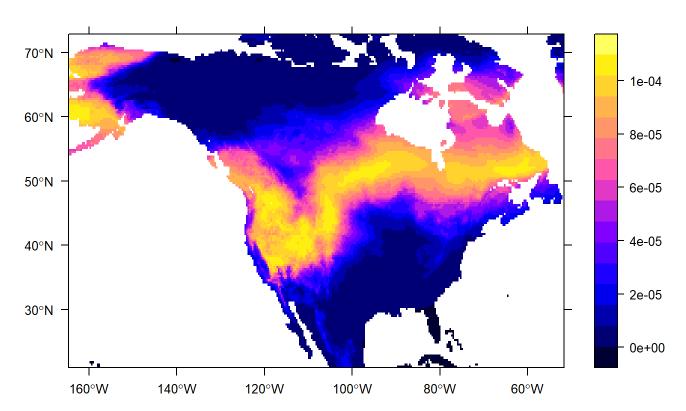
# **Probability Density Surface for b**



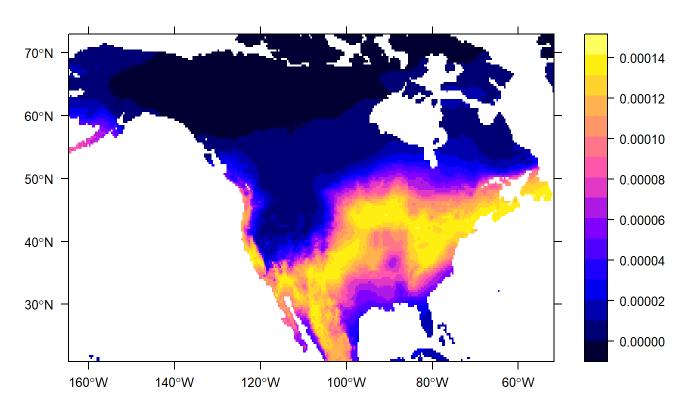
# **Probability Density Surface for c**



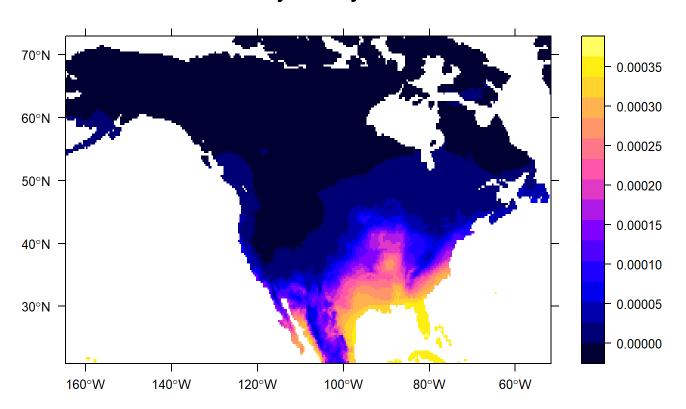
# Probability Density Surface for d



# Probability Density Surface for e

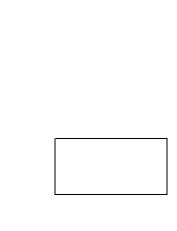


#### **Probability Density Surface for f**



# **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)
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 <- 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

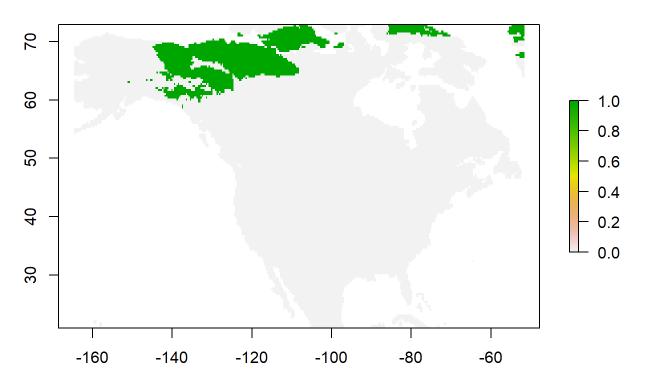
```
oddsRatio(asn, pp12)
## $`P1/P2 odds ratio`
           а
                                      С
## 1.146839e-08 8.417012e-06 4.205954e-03 1.431251e+00 3.316179e+02
## 5.230446e+04
## $`odds of a pixel to the odds of the max/min pixel`
## ratioToMax.a ratioToMax.b ratioToMax.c ratioToMax.d ratioToMax.e
## 1 1.626156e-11 2.111854e-06 0.002773832 0.02499529 1.166192978
## 2 1.080405e-02 2.359802e-01 0.257425866 0.13306854 0.003307999
## ratioToMax.f ratioToMin.a ratioToMin.b ratioToMin.c ratioToMin.d
## 1 1.241220e-01 6.034369e+10 313.3168 1.113673e+04 9.275299e+19
## 2 9.263921e-07 1.793020e+12 1439.8675 3.025310e+11 2.208380e+13
## ratioToMin.e ratioToMin.f
## 1 1.730175e+08 498340.6
## 2 2.018424e+01
                    1088710.9
```

# **Binary reclassification**

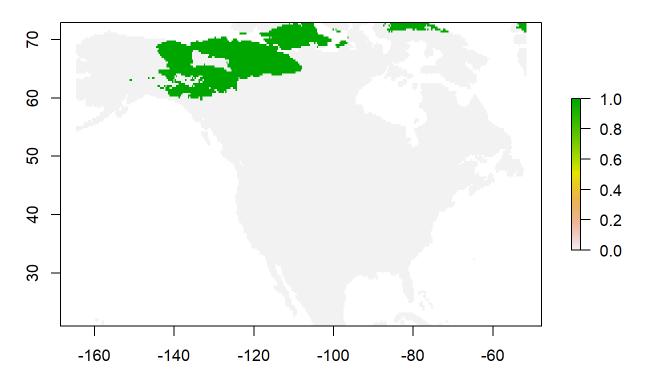
### Top 10% of probability surface (defined by % area)

```
qtlRaster(asn, threshold = 0.1, thresholdType = 2)
```

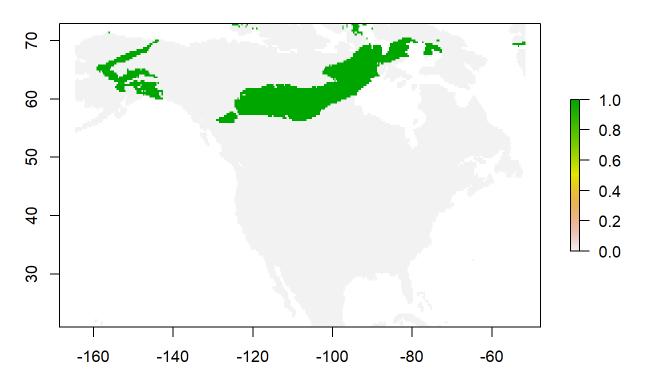
Top 10% by Area for a



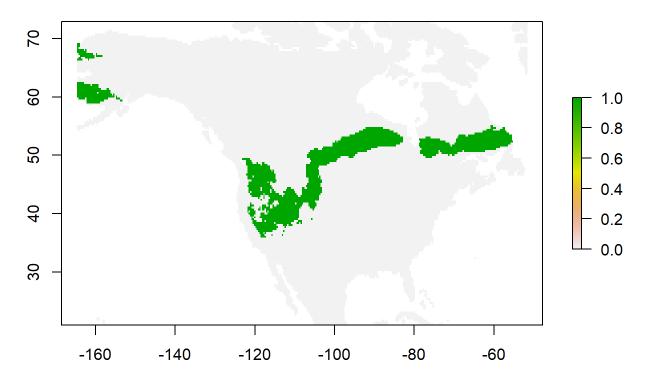
Top 10% by Area for b



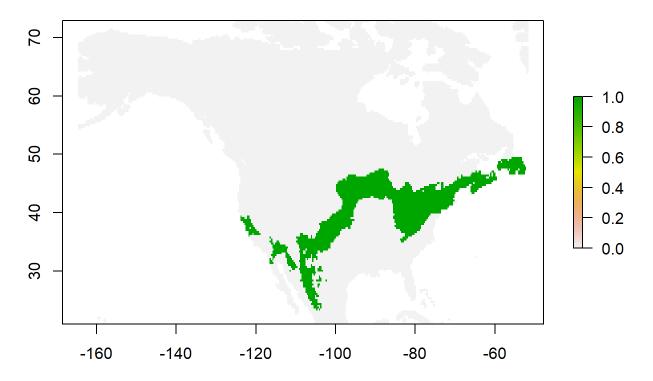
Top 10% by Area for c



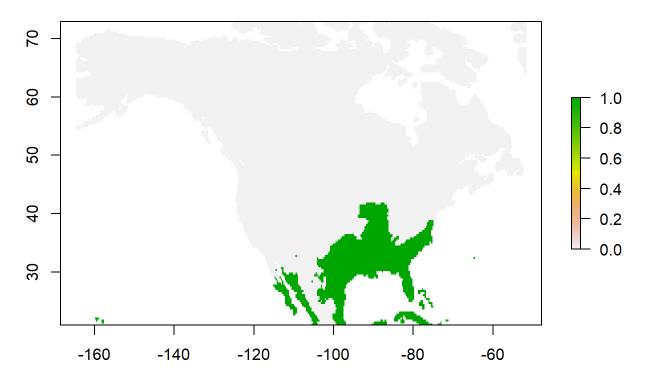
Top 10% by Area for d



Top 10% by Area for e



#### Top 10% by Area for f

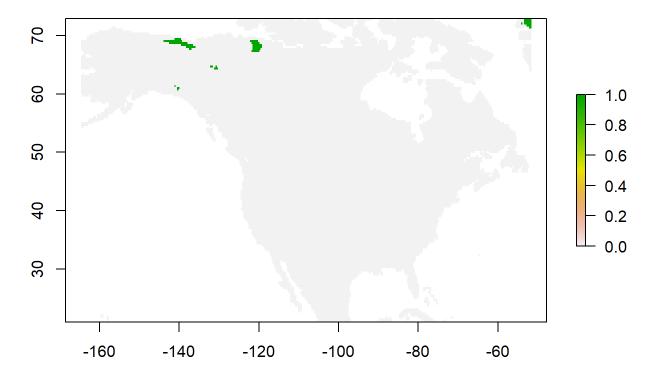


```
## class : RasterStack
## dimensions : 156, 339, 52884, 6 (nrow, ncol, ncell, nlayers)
## resolution : 0.3333332, 0.3333332 (x, y)
## extent : -164.6667, -51.66672, 20.91662, 72.9166 (xmin, xmax, ymin, ymax)
## crs : +proj=longlat +datum=WGS84 +no_defs +ellps=WGS84
+towgs84=0,0,0
## names : a, b, c, d, e, f
## min values : 0, 0, 0, 0, 0, 0
## max values : 1, 1, 1, 1, 1
```

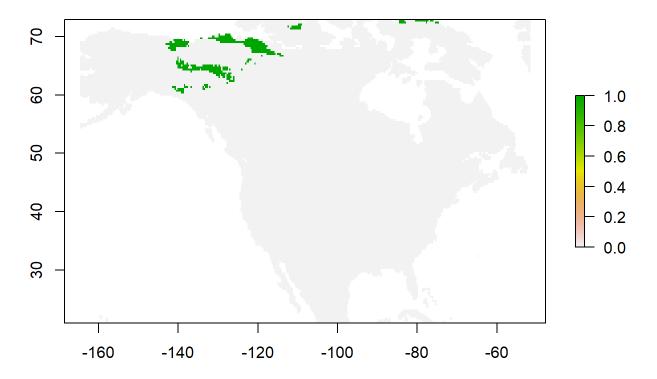
# Top 10% of probability surface (defined by % cumulative probability)

```
qtlRaster(asn, threshold = 0.1, thresholdType = 1)
```

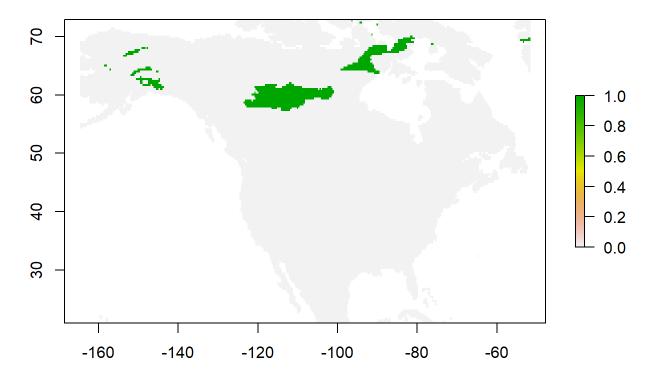
Top 10% by Cumulative Probability for a



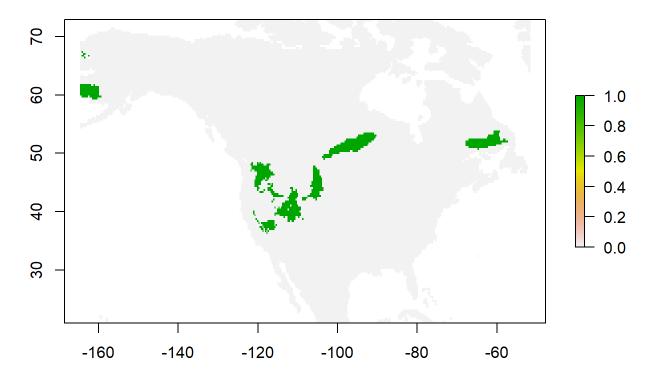
Top 10% by Cumulative Probability for b



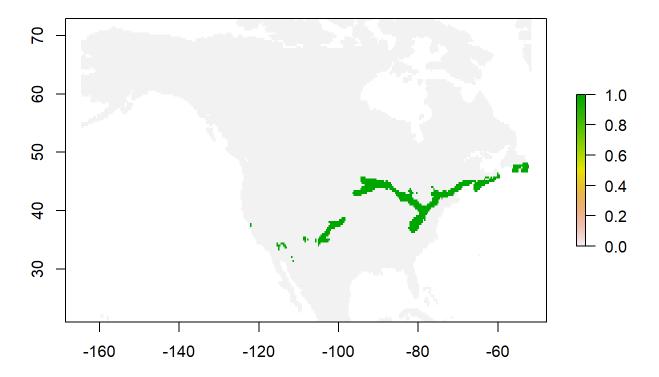
Top 10% by Cumulative Probability for c



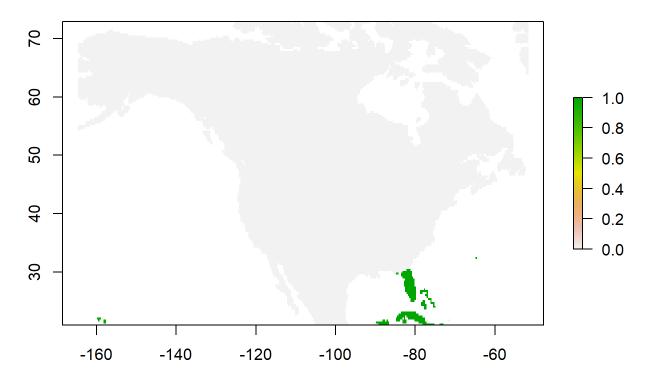
Top 10% by Cumulative Probability for d



Top 10% by Cumulative Probability for e



Top 10% by Cumulative Probability for f

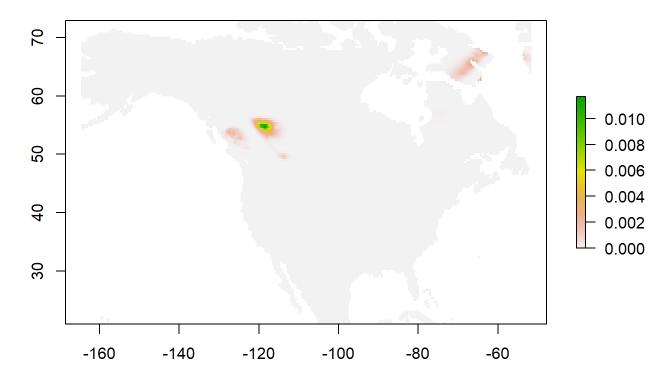


```
## class : RasterStack
## dimensions : 156, 339, 52884, 6 (nrow, ncol, ncell, nlayers)
## resolution : 0.3333332, 0.3333332 (x, y)
## extent : -164.6667, -51.66672, 20.91662, 72.9166 (xmin, xmax, ymin, ymax)
## crs : +proj=longlat +datum=WGS84 +no_defs +ellps=WGS84
+towgs84=0,0,0
## names : a, b, c, d, e, f
## min values : 0, 0, 0, 0, 0, 0
## max values : 1, 1, 1, 1, 1
```

# Joint probability for individuals of common origin

jointP(asn)

#### **Joint Probability**

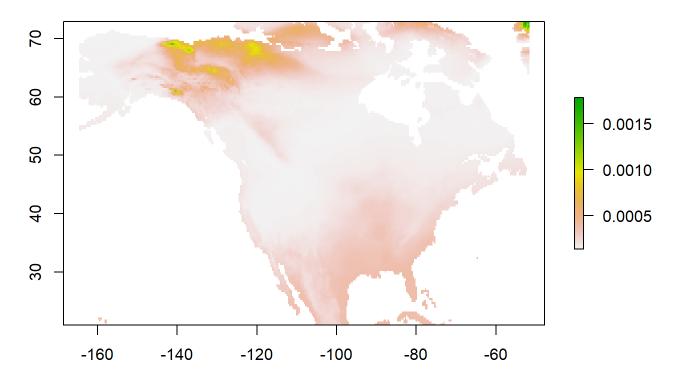


```
: RasterLayer
## dimensions : 156, 339, 52884 (nrow, ncol, ncell)
## resolution : 0.3333332, 0.3333332 (x, y)
            : -164.6667, -51.66672, 20.91662, 72.9166 (xmin, xmax,
## extent
ymin, ymax)
             : +proj=longlat +datum=WGS84 +no defs +ellps=WGS84
## crs
+towgs84=0,0,0
## source
             : memory
             : Joint_Probability
## names
             : 1.031402e-44, 0.01170466
                                          (min, max)
## values
```

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

unionP(asn)

#### **Union Probability**

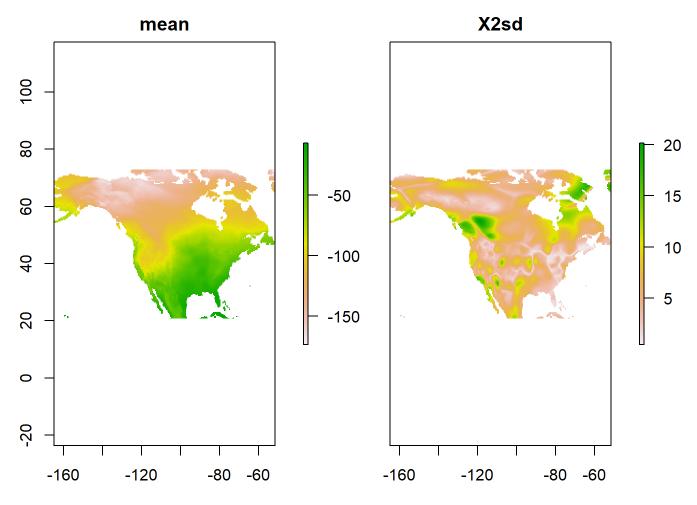


```
: RasterLayer
## dimensions : 156, 339, 52884 (nrow, ncol, ncell)
## resolution : 0.3333332, 0.3333332 (x, y)
             : -164.6667, -51.66672, 20.91662, 72.9166 (xmin, xmax,
## extent
ymin, ymax)
              : +proj=longlat +datum=WGS84 +no defs +ellps=WGS84
## crs
+towgs84=0,0,0
## source
             : memory
## names
              : layer
              : 0.0001361405, 0.001787688
## values
                                           (min, max)
```

# Quality analysis of geographic assignment

```
# oxygen and hydrogen isotopes of known-origin bird
data(bird_isotope)

# crop the world hydrogen data to North America
r <- crop(d2h_world, naMap)
plot(r)</pre>
```



# convert 2 standard deviation from d2h\_world to 1 standard deviation r[[2]] <- r[[2]]/2

```
# seperate the hydrogen isotope for the known-origin bird
bird_d2h <- bird_isotope[1:20,c("Longitude", "Latitude", "d2H")]
coordinates(bird_d2h) <- c(1,2)
proj4string(bird_d2h) <- proj4string(d2h_world)</pre>
```

 $\mbox{\#}$  run quality assessment based hydrogen isotope from precipitation and known-origin bird



