Operations on Raster Data I

HES 505 Fall 2023: Session 13

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Objectives

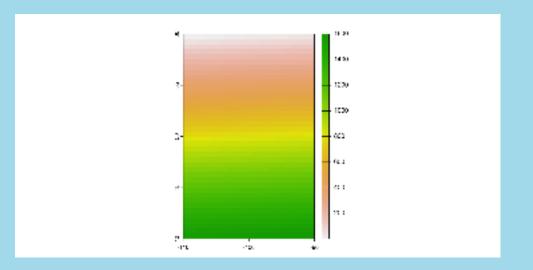
- By the end of today, you should be able to:
 - Align rasters for spatial processing
 - Adjust the resolution of raster data
 - Combine (or reduce) rasters to match the extent of your analysis

Aligning rasters for spatial processing

Projecting raster data

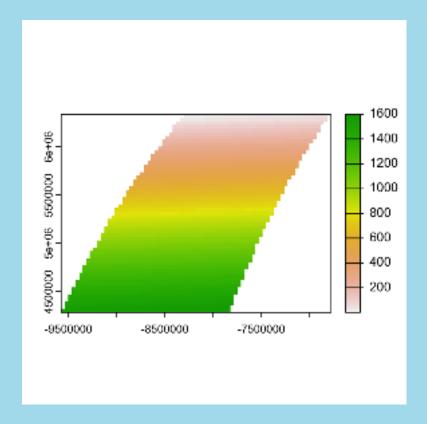
- Transformation from lat/long to planar CRS involves some loss of precision
- New cell values estimated using overlap with original cells
- Interpolation for continuous data, nearest neighbor for categorical data
- Equal-area projections are preferred;
 especially for large areas

```
library(sf)
library(terra)
library(spDataLarge)
r <- rast(xmin=-110, xmax=-90, ymin=40, ymax=60, not values(r) <- 1:ncell(r)
plot(r)</pre>
```

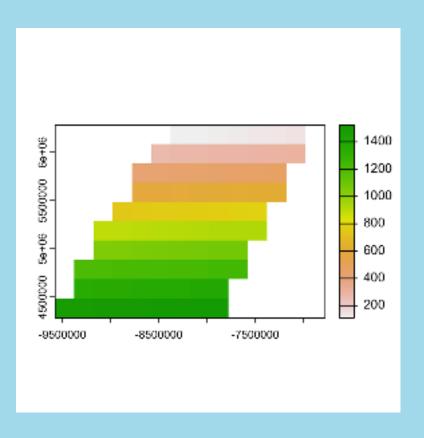


Projecting raster data

- simple method; alignment providing a template to not guaranteed
 - newcrs <- "+proj=robin +datum=WGS8</pre>
 - pr1 <- terra::project(r, newcrs)</pre>
 - plot(pr1)

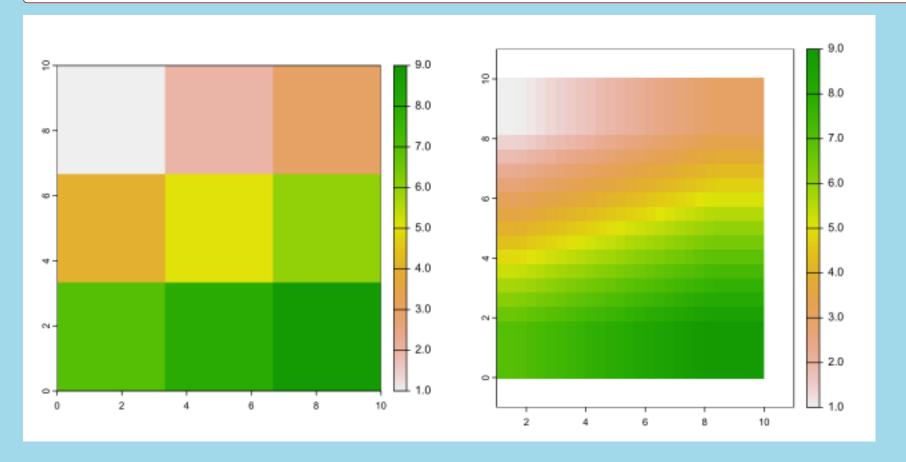


ensure alignment



Aligning Data: resample

```
1 r <- rast(nrow=3, ncol=3, xmin=0, xmax=10, ymin=0, ymax=10)
2 values(r) <- 1:ncell(r)
3 s <- rast(nrow=25, ncol=30, xmin=1, xmax=11, ymin=-1, ymax=11)
4 x <- resample(r, s, method="bilinear")</pre>
```



Adjusting resolution

Downscaling and Upscaling

- Aligning data for later analysis
- Remembering scale
- Thinking about support

Changing resolutions

- aggregate, disaggregate, resample allow changes in cell size
- aggregate requires a function (e.g., mean() or min()) to determine what to do with the grouped values
- resample allows changes in cell size and shifting of cell centers (slower)

Changing resolutions: aggregate

1 r <- rast() 2 r

class : SpatRaster

dimensions : 180, 360, 1 (nrow,

ncol, nlyr)

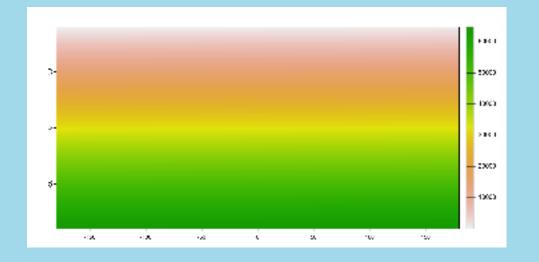
resolution : 1, 1 (x, y)

extent : -180, 180, -90, 90

(xmin, xmax, ymin, ymax)

coord. ref. : lon/lat WGS 84

- 1 values(r) <- 1:ncell(r)</pre>
- 2 plot(r)



```
1 ra <- aggregate(r, 20)
2 ra</pre>
```

class : SpatRaster

dimensions: 9, 18, 1 (nrow, ncol,

nlyr)

resolution : 20, 20 (x, y)

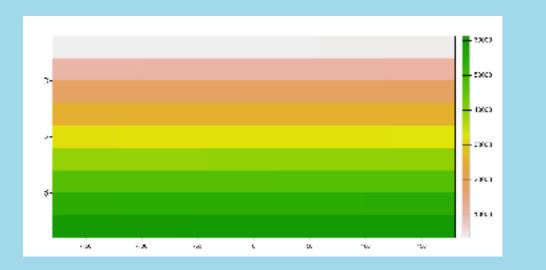
extent : -180, 180, -90, 90

(xmin, xmax, ymin, ymax)

coord. ref. : lon/lat WGS 84

source(s) : memory
name : lyr.1
min value : 3430.5
max value : 61370.5

1 plot(ra)



Changing resolutions: disagg

```
1 ra <- aggregate(r, 20)
2 plot(ra)</pre>
```

```
5-1003
-1003
-1003
-1003
-1003
-1003
```

```
1 rd \leftarrow disagg(r, 20)
 1 rd
class : SpatRaster
dimensions : 3600, 7200, 1 (nrow,
ncol, nlyr)
resolution : 0.05, 0.05 (x, y)
extent : -180, 180, -90, 90
(xmin, xmax, ymin, ymax)
coord. ref.: lon/lat WGS 84
source :
spat 4QyAlALziT1WmUe 97640.tif
name
           : lyr.1
min value : 1
max value : 64800
```

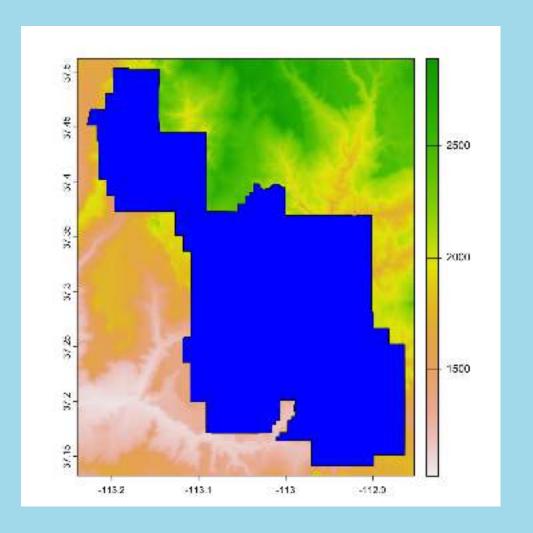
```
1 plot(rd)
```



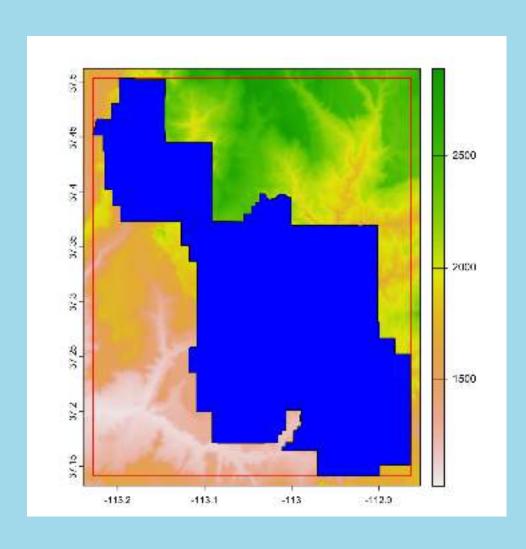
Modifying the Extent

Dealing with Different Extents

- Raster extents often larger than our analysis
- Reducing memory and computational resources
- Making attractive maps



Using terra::crop()



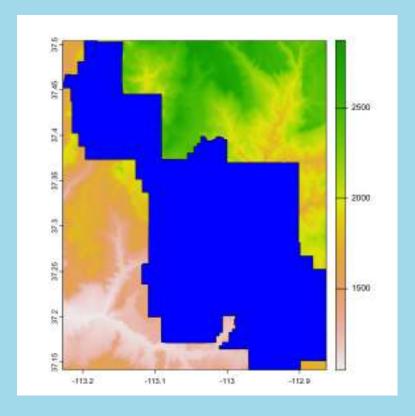
- Coordinate Reference System must be the same for both objects
- Crop is based on the (converted)
 SpatExtent of the 2nd object
- **snap** describes how **y** will be aligned to the raster
- Returns all data within the extent

Using terra::crop()

```
library(sf)
library(terra)
library(spDataLarge)
srtm = rast(system.file("raster/srtm.tif", package = "spDataLation = read_sf(system.file("vector/zion.gpkg", package = "spDataLation = st_transform(zion, crs(srtm))
crs(srtm) == crs(zion)
```

[1] TRUE

```
1 srtm.crop <- crop(x=srtm, y=zion, snap="near")</pre>
```

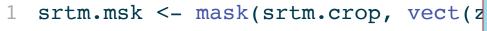


Using mask()

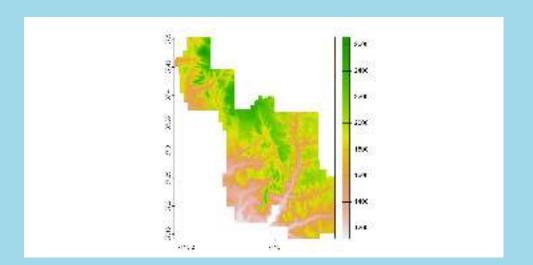
- Often want to get rid of all values outside of vector
- Can set mask=TRUE in crop() (y must be SpatVector)
- Or use mask()

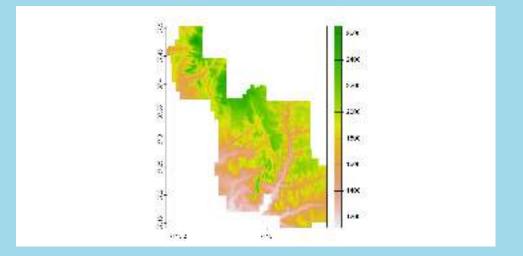
```
1 srtm.crop.msk <- crop(x=srtm, y=ve
```

2 plot(srtm.crop.msk)



2 plot(srtm.msk)





Using mask()

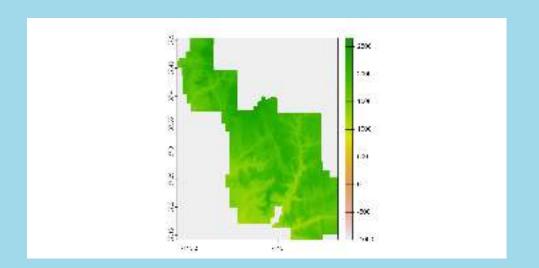
- Allows more control over what the mask does
- Can set maskvalues and updatevalues to change the resulting raster
- Can also use **inverse** to mask out the vector

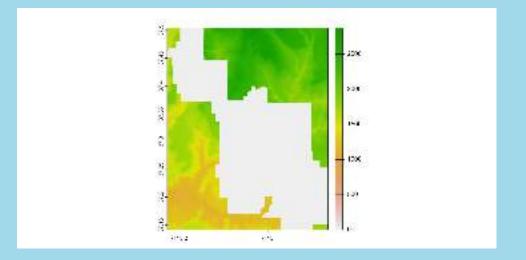
```
1 srtm.msk <- mask(srtm.crop, vect(z</pre>
```

2 plot(srtm.msk)



2 plot(srtm.msk)





Extending boundaries

- Vector slightly larger than raster
- Especially when using buffered datasets
- Can use extend
- Not exact; depends on snap()

```
1 zion.buff <- zion %>%
2 st_buffer(., 10000)
3 srtm.ext <- extend(srtm, vect(zion ext(srtm.ext))</pre>
```

SpatExtent : -113.343749879444,
-112.74791654615, 37.0479167631968,
37.5979167631601 (xmin, xmax, ymin,
ymax)

1 ext(vect(zion.buff))

SpatExtent : -113.343652923976,
-112.747986193365, 37.0477357596604,
37.5977812137969 (xmin, xmax, ymin,
ymax)

