

# Rediscretizing Tracks

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**These analyses only consider the first two releases, intended as Phase 1 (phase 2 = survival study)**

However, there were 221 additional fish released in phases 3 - 5 that could be incorporated. They were released at higher flows, which would add interesting complexity, and additional analysis needs.

## Rediscretization of Tracks

- Using primary and secondary filtered data to rediscretize tracks for further analysis
- Tracks have been split into bursts where successive positions were separated by  $> 50\text{m}$
- this threshold can be altered in “Final\_Filtering.Rmd” if desired
- Before rediscretizing, remove bursts with  $< 10$  positions (too few to rediscretize in adehabitatLT)
- After, recalculated migration speed between positions

```
load("Maestros/AllFish_FiltSec4Bursts.RData") # single object, named red7
dim(red7) # 81895 detections
```

```
## [1] 81895    16
```

```
length(unique(red7$id)) # 430 unique fish
```

```
## [1] 430
```

```
length(unique(red7$burst)) # 665 unique bursts
```

```
## [1] 665
```

```
ndetects.fish = summarize(group_by(red7, id), ndet = n())
mean(ndetects.fish$ndet) # 190.5 per fish
```

```
## [1] 190.4535
```

```
range(ndetects.fish$ndet) # ranges from 17 - 578
```

```
## [1] 17 578
```

```
ndetects.burst = summarize(group_by(red7, burst), ndet = n())
mean(ndetects.burst$ndet) # 123.2 per burst
```

```
## [1] 123.1504
```

```
range(ndetects.burst$ndet) # ranges from 3 - 578
```

```
## [1] 3 578
```

```
max(red7$spd_mps, na.rm=T) # 42.8 mps
```

```
## [1] 42.82635
```

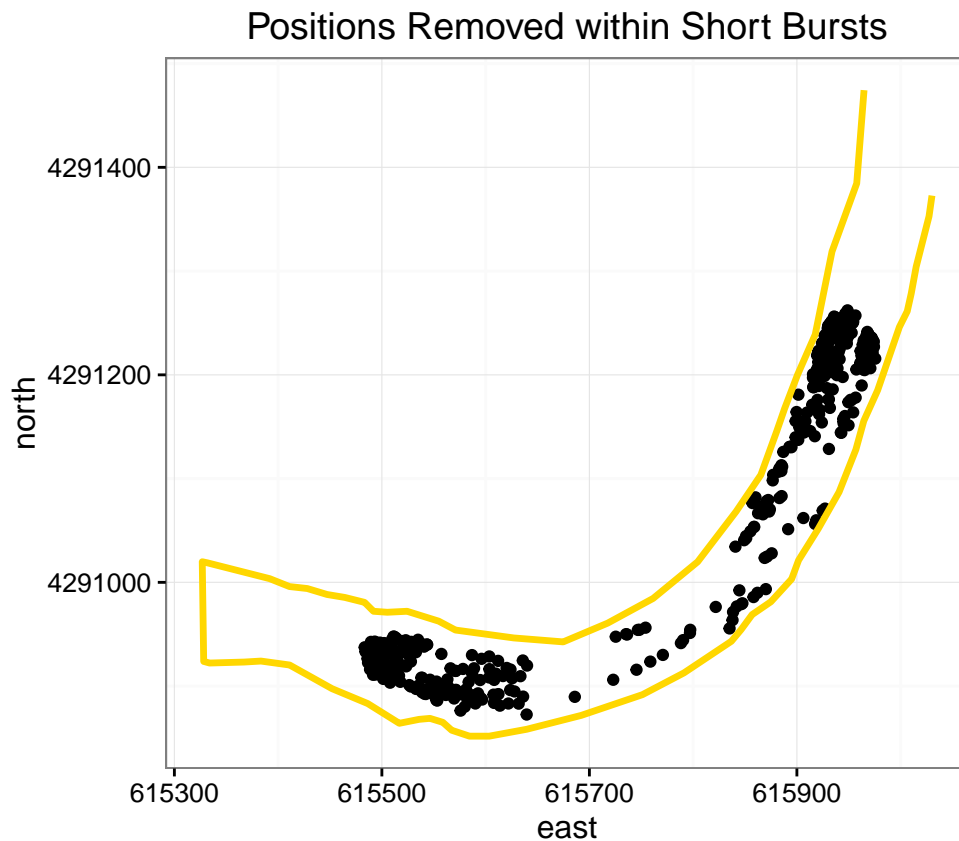
```
bursts.rem = data.frame(ndetects.burst[ndetects.burst$ndet<10,])  
nrow(bursts.rem) # 104 bursts removed
```

```
## [1] 107
```

```
sum(bursts.rem$ndet) # 535 positions removed + 166 removed when cut the bursts (<4 pos remaining in
```

```
## [1] 535
```

```
red.br = red7[(red7$burst %in% bursts.rem$burst),]  
  
river3 = fortify(river3)  
ggplot(data = red.br, aes(x=east, y=north)) + geom_point() +  
  geom_path(data = river3, aes(long, lat), col="gold", size=1.2 ) +  
  ggtitle(label = "Positions Removed within Short Bursts") +  
  theme_bw() + coord_fixed()
```



```

red8 = red7[!(red7$burst %in% bursts.rem$burst),]

# prep object for redisltraj()
red8.ltraj = as.ltraj(xy=red8[,c("east","north")], date=red8$date,
                     id=red8$id, burst = factor(red8$burst),
                     infolocs=red8[,c("Hpes","east","north")])

# discretize in space
redlt.ssrz = (redisltraj(red8.ltraj, u=27, type="space", nnew=5))
# u=25 is the smallest distance where the code will work.

# convert back to dataframe
red8.ssrz = ld(redlt.ssrz)
red8.ssrz=red8.ssrz[order(red8.ssrz$id,red8.ssrz$date),]

# recalculate migration speed
red8.ssrz$spd_mps = red8.ssrz$dist / red8.ssrz$dt

dim(red8.ssrz) # 9040 detections after discretization

## [1] 9040 15

length(unique(red8.ssrz$id)) # 430

## [1] 430

ndetects.discr = summarize(group_by(red8.ssrz, id), ndet = n())
mean(ndetects.discr$ndet) # 21.0 per fish

## [1] 21.02326

range(ndetects.discr$ndet) # ranges from 5-27

## [1] 5 27

max(red8.ssrz$spd_mps, na.rm=T) # 20.5 mps

## [1] 20.49389

```

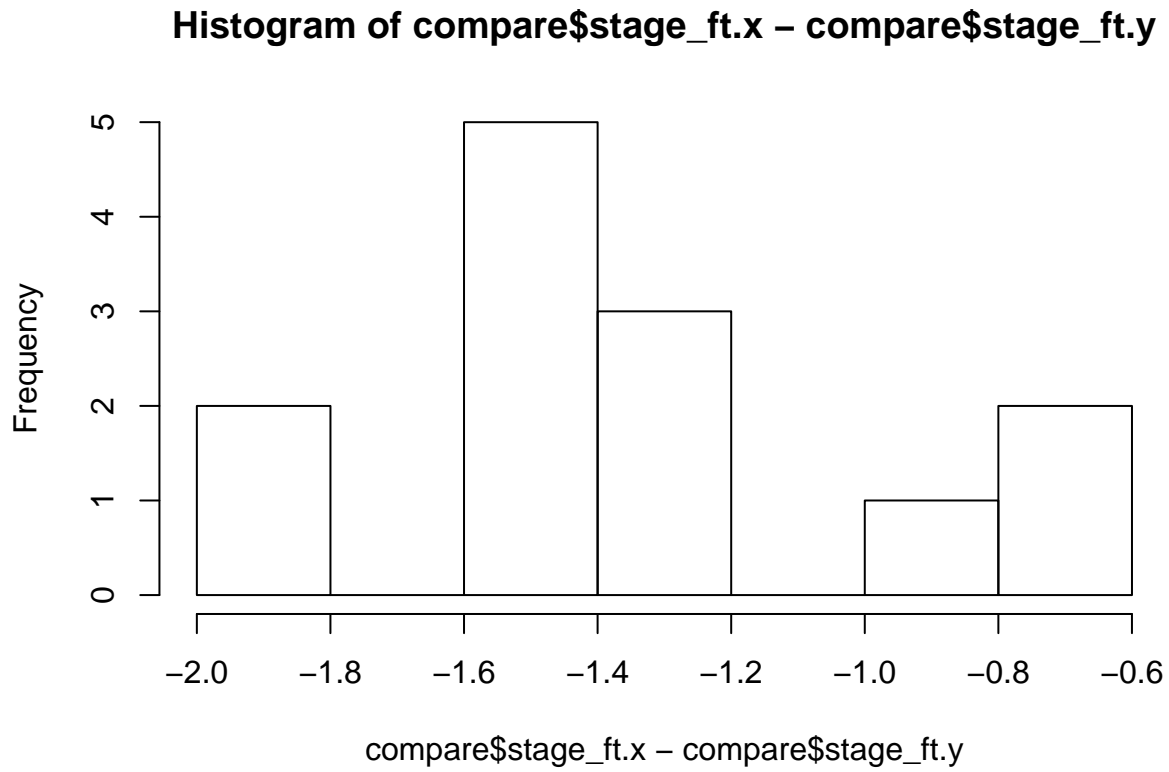
## Add covariates (release event, river stage) back into dataframe to possibly use later

\*Compared un-QAQCed data on river stage from CDEC station (no discharge available) with USGS measured values

Save the discretized object to RData

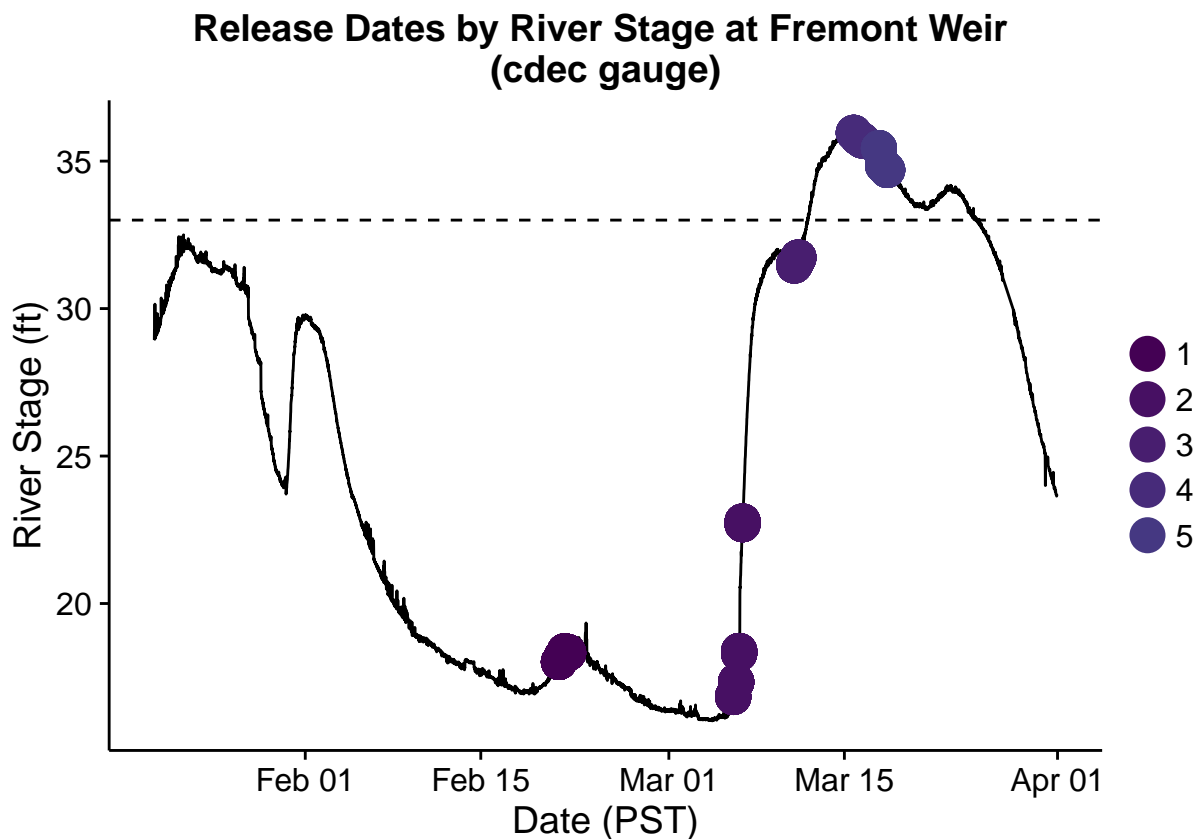
Note: In 2015 I removed tracks which had < 10 positions after spatial discretization. However, this year's analysis is slightly different since I'm working with bursts -> we should have already controlled for sparse tracks (i.e., those where interpolation might be inappropriate). Plus, above I've removed any burst with < 10 positions. Therefore, I'm imposing no additional filtering after discretization.

```
## [1] -1.27 -1.26 -1.28 -0.85 -0.79 -1.84 -1.88 -1.58 -1.56 -1.41 -1.43
## [12] -1.45 -0.71    NA
```



```
## [1] -1.331538
```

```
## [1] 0.3681224
```



And finally, output the general metrics about the remaining dataset

```
dim(red8.ssrdez) # 9040 detections after discretization
```

```
## [1] 9040 31
```

```
length(unique(red8.ssrdez$id)) # 430
```

```
## [1] 430
```

```
ndetects.discr = summarize(group_by(red8.ssrdez, id), ndet = n())
mean(ndetects.discr$ndet) # 21.0 per fish
```

```
## [1] 21.02326
```

```
range(ndetects.discr$ndet) # ranges from 5 - 27
```

```
## [1] 5 27
```

```
max(red8.ssrdez$spd_mps, na.rm=T) # 20.49 mps
```

```
## [1] 20.49389
```