

# Metrics for Recorder behaviour

*Tom August*

*10 October 2016*

## Metrics

We are going to split metric into three broad groups: *Engagement profile*, *Spatial*, and *Taxanomic*

## Temporal Metrics

These metrics measure the recording pattern across time such as the number of days that a recorder produces records. These have been termed engagement profiles by others., The metrics here are from Ponciano and Brasileiro 2014 who used the metrics on participant of zooniverse projects. The metrics were also used by Boakes *et al* 2016.

## Summer period

One issue that we have across these metrics and some others is that recording is not consistent across the year and so there can be issues with the numbers generated. To address this the data can be subset to only the summer period, when recorders are active. This period needs to be defined in such a way that the same method can be used across taxanomic groups and will be robust to changes in the start and end of the summer period from year to year.

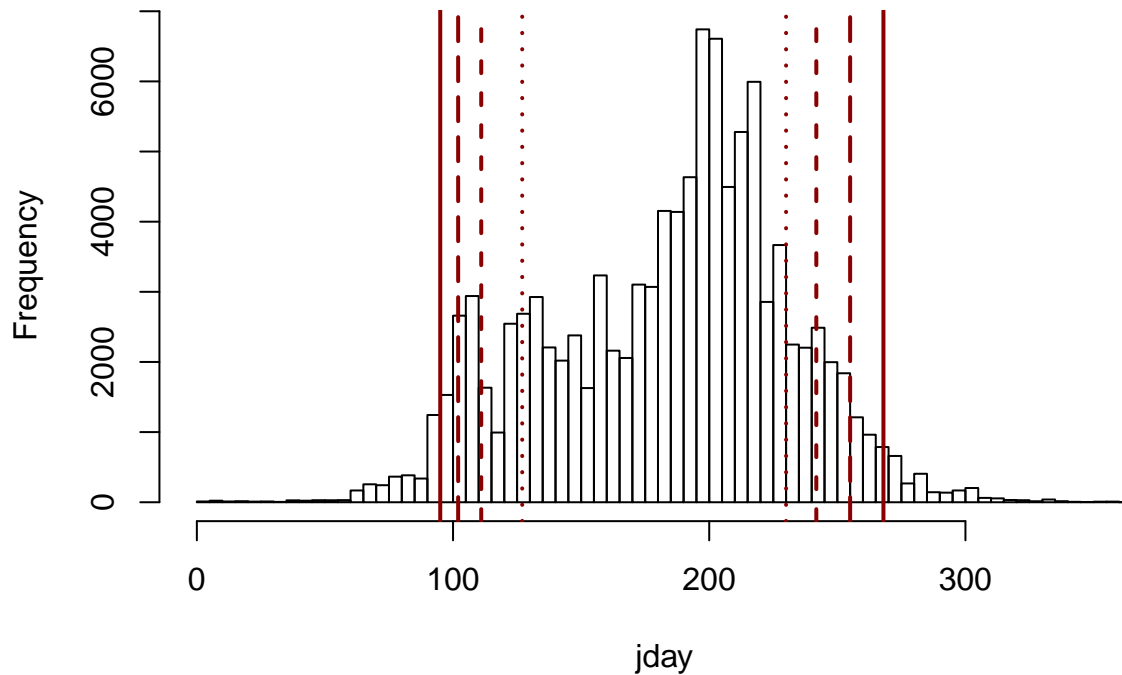
I suggest we use a percentage cut off, for example take the period of the year that contains 90% of the data. Lets have a look at how this might work

```
# First lets have a look at the distribution of
# records through the year
jday <- as.POSIXlt(iRB$date_start)$yday

# Where are the limits of different percentiles
p95 <- quantile(jday, c(0.025,0.975))
p90 <- quantile(jday, c(0.05,0.95))
p80 <- quantile(jday, c(0.1,0.9))
p70 <- quantile(jday, c(0.15,0.85))

hist(jday, 100, main = paste('Histogram of recording day with\n',
                             'cutoffs at 95%, 90%, 80%, & 70%'))
abline(v = p95, lty = 1, col = 'darkred', lwd = 2)
abline(v = p90, lty = 5, col = 'darkred', lwd = 2)
abline(v = p80, lty = 2, col = 'darkred', lwd = 2)
abline(v = p70, lty = 3, col = 'darkred', lwd = 2)
```

## Histogram of recording day with cutoffs at 95%, 90%, 80%, & 70%



It looks like 90% might be a good value to go for in this case. We would then need a function that could create these values for each year of the data and throw out data that was outside the summer periods

```
summerData <- function(data, probs = c(0.05, 0.95),
                        date_col = 'date_start'){

  # check date column
  if(!inherits(data[, date_col], 'Date')){
    stop('Your date column is not a date')
  }

  # create J-day column
  data$Jday <- as.POSIXlt(data[,date_col])$yday

  # create year column
  data$year <- as.POSIXlt(data[,date_col])$year+1900

  # create summer column
  data$summer <- FALSE

  qsf <- qsl <- NULL

  # now for each year loop through and create an
  # index column
  for(i in sort(unique(data$year))){
```

```

year_quantiles <- quantile(data$Jday[data$year == i], probs = probs)
qsf <- c(qsf, year_quantiles[1])
qsl <- c(qsl, year_quantiles[2])
data$summer[data$Jday >= year_quantiles[1]
            & data$Jday <= year_quantiles[2]
            & data$year == i] <- TRUE
}

summer_data <- data[data$summer, ]
attr(summer_data, 'cutoffs') <- data.frame(year = sort(unique(data$year)),
                                           quantile_first = qsf,
                                           quantile_last = qsl)

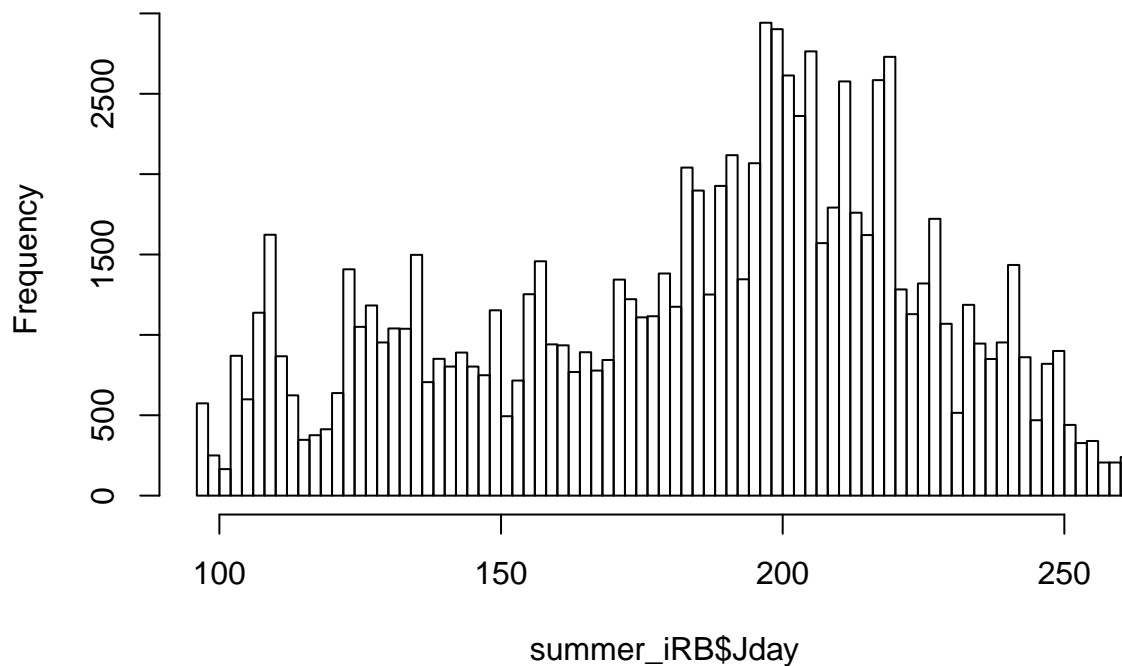
return(summer_data)
}

summer_iRB <- summerData(data = iRB,
                        probs = c(0.05, 0.95),
                        date_col = 'date_start')

# Look at the 'cut' data
hist(summer_iRB$Jday, 100)

```

**Histogram of summer\_iRB\$Jday**



```

# Here are the cuts
attr(summer_iRB, 'cutoffs')

```

```
##   year quantile_first quantile_last
## 1 2014           104           260
## 2 2015           97           262
## 3 2016          107           250
```

## Activity ratio

*“The proportion of days on which the volunteer was active in relation to the total days he/she remained linked to the project”* (Ponciano and Brasileiro 2014)

```
# Create a function to calculate activity ratio
activityRatio <- function(recorder_name,
                          data,
                          recorder_col = 'recorders',
                          date_col = 'date_start'){

  # check date column
  if(!inherits(data[, date_col], 'Date')){
    stop('Your date column is not a date')
  }

  # Get the recorders data
  data <- data[data[, recorder_col] == recorder_name, ]

  # Get unique dates
  dates <- unique(data[, date_col])

  # Get the first and last date
  first_last <- range(dates)

  # Total duration of this recorder
  duration <- as.numeric(first_last[2] - first_last[1]) + 1

  # calculate ratio
  activity_ratio <- length(dates)/duration

  # return
  return(data.frame(recorder = recorder_name,
                    activity_ratio = activity_ratio,
                    total_duration = duration,
                    active_days = length(dates)))
}

# Test on David and Tom
activityRatio(data = summer_iRB, recorder_name = 'Roy, David')
```

```
##   recorder activity_ratio total_duration active_days
## 1 Roy, David    0.1509217           868          131
```

```
activityRatio(data = summer_iRB, recorder_name = 'August, Tom')
```

```
##   recorder activity_ratio total_duration active_days
```

```
## 1 August, Tom      0.01587302      441      7
```

```
## David is a more active recorder than Tom ##
```

```
# Run for everyone
```

```
all_AR <- do.call(rbind, lapply(X = unique(iRB$recorders),  
                                FUN = activityRatio,  
                                data = summer_iRB))
```

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## Warning in min.default(structure(numeric(0), class = "Date"), na.rm =
## FALSE): no non-missing arguments to min; returning Inf

## Warning in max.default(structure(numeric(0), class = "Date"), na.rm =
## FALSE): no non-missing arguments to max; returning -Inf

## Warning in min.default(structure(numeric(0), class = "Date"), na.rm =
## FALSE): no non-missing arguments to min; returning Inf

## Warning in max.default(structure(numeric(0), class = "Date"), na.rm =
## FALSE): no non-missing arguments to max; returning -Inf

## Warning in min.default(structure(numeric(0), class = "Date"), na.rm =
## FALSE): no non-missing arguments to min; returning Inf

## Warning in max.default(structure(numeric(0), class = "Date"), na.rm =
## FALSE): no non-missing arguments to max; returning -Inf

## Warning in min.default(structure(numeric(0), class = "Date"), na.rm =
## FALSE): no non-missing arguments to min; returning Inf

## Warning in max.default(structure(numeric(0), class = "Date"), na.rm =
## FALSE): no non-missing arguments to max; returning -Inf
```

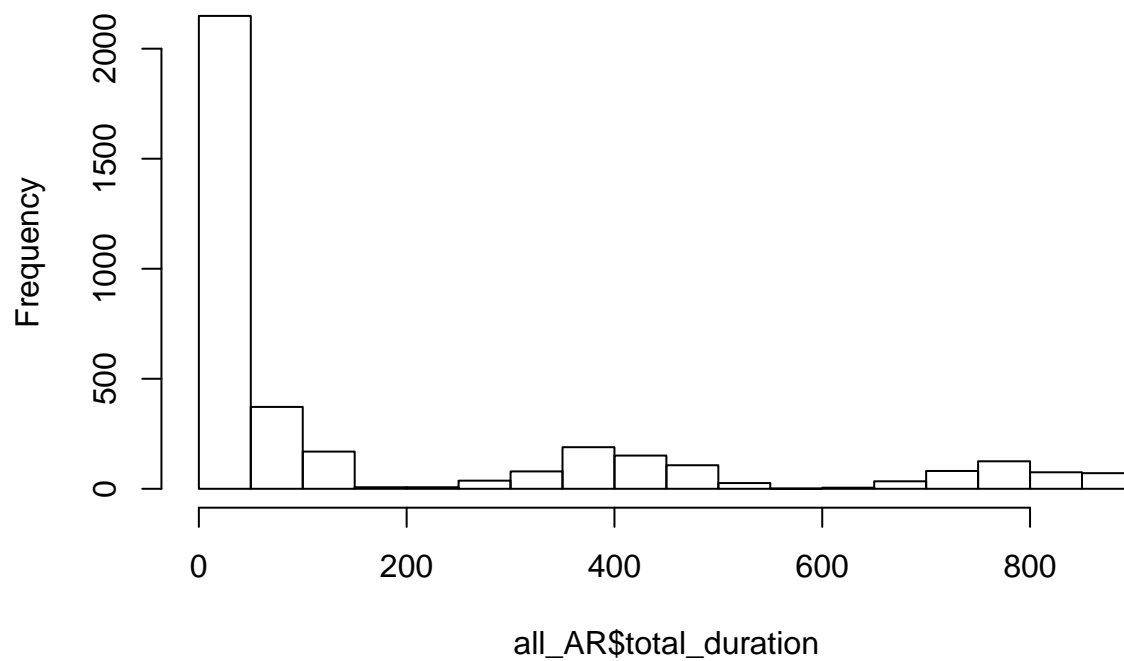
```
# Lets have a look at some of these
head(all_AR, 20)
```

	recorder	activity_ratio	total_duration	active_days
## 1	Brookes , Anne	0.07168894	823	59
## 2	Burgoyne, Steve	0.39830508	118	47
## 3	Brown, Peter	0.11309524	504	57
## 4	Rutherford, Joanna	0.01292597	851	11
## 5	Allan, David	0.29532497	877	259
## 6	Millward, Martin	0.08000000	50	4
## 7	Foulkes-Arellano, Paul	0.02192448	821	18
## 8	Stewart, Tam	0.26339795	877	231
## 9	Forbes, Andrew	0.20833333	24	5
## 10	Richardson, Rosie	0.28000000	25	7
## 11	Partridge, Francesca	0.18002466	811	146
## 12	Card , Graeme	0.05378973	409	22
## 13	Honey, Hawk	0.08421053	855	72
## 14	Melzack, David	0.31250000	96	30
## 15	Povall, Ed	0.10000000	30	3
## 16	Goodwin, Paul	0.15700483	414	65
## 17	Coulson, Joe	0.18965517	116	22
## 18	Bailey, Peggy	0.33064516	124	41
## 19	Roy, David	0.15092166	868	131
## 20	Woodley, Caroline	0.14634146	123	18

I think this metric tells a story in a combination of the ratio and the total number of days. I think the ratio means more when the recorder has been recording for a long duration

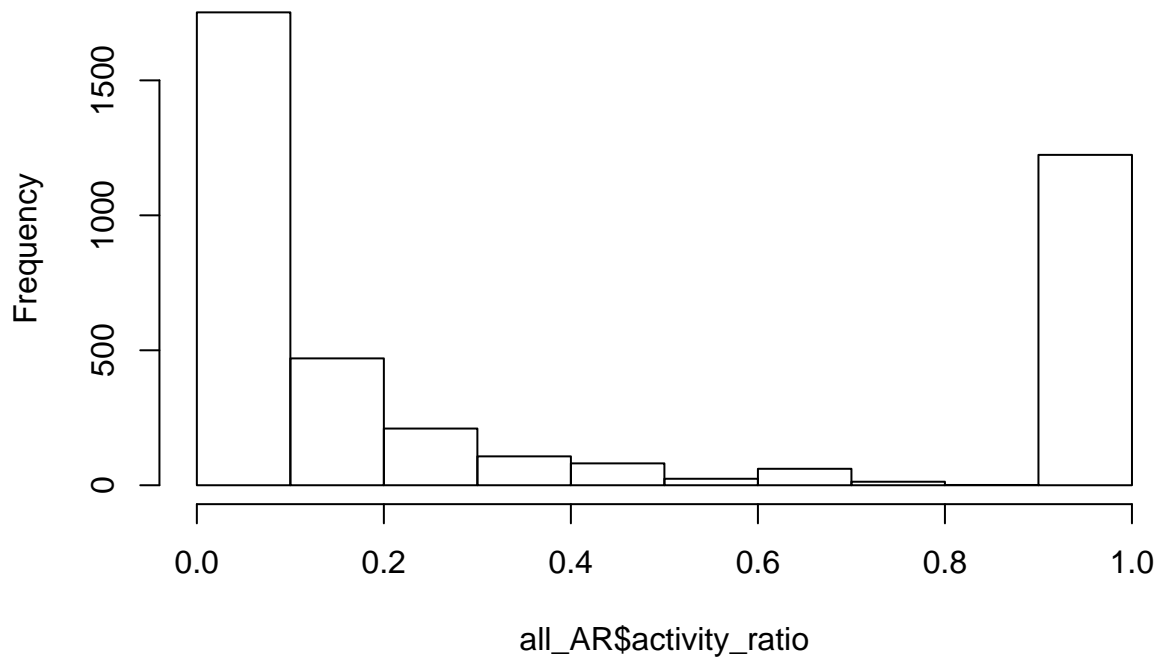
```
# Have a look at the distribution of these 2 metrics  
# There looks like there could be an effect of year  
hist(all_AR$total_duration, breaks = 30)
```

### Histogram of all\_AR\$total\_duration



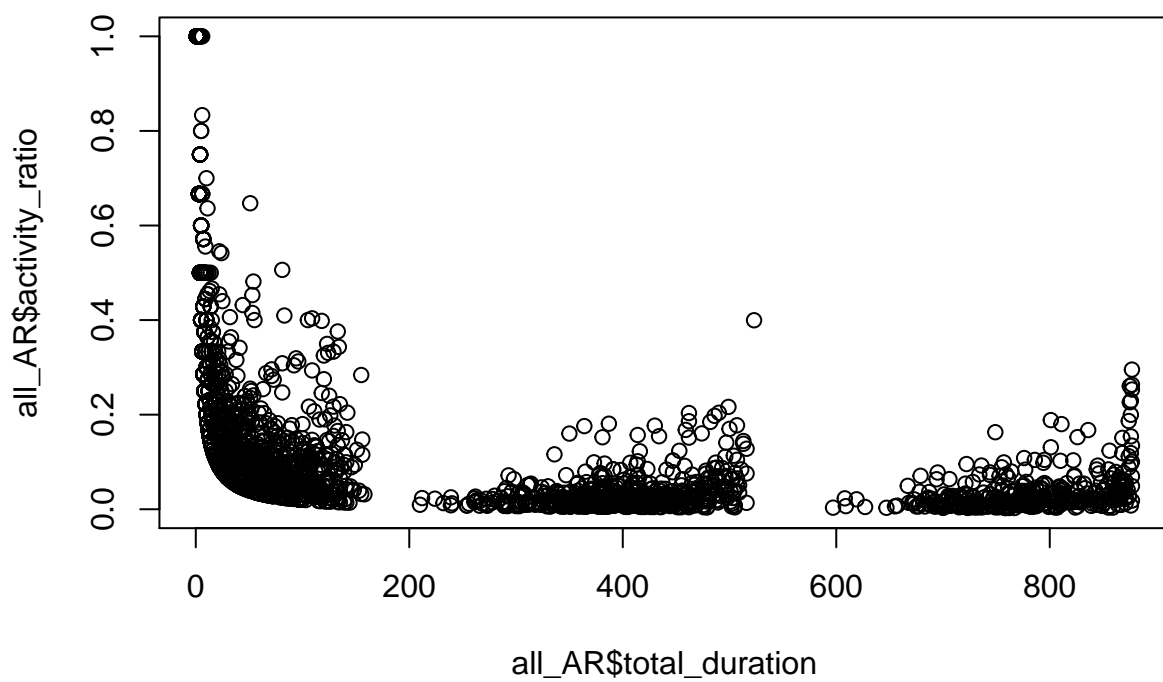
```
hist(all_AR$activity_ratio)
```

## Histogram of all\_AR\$activity\_ratio

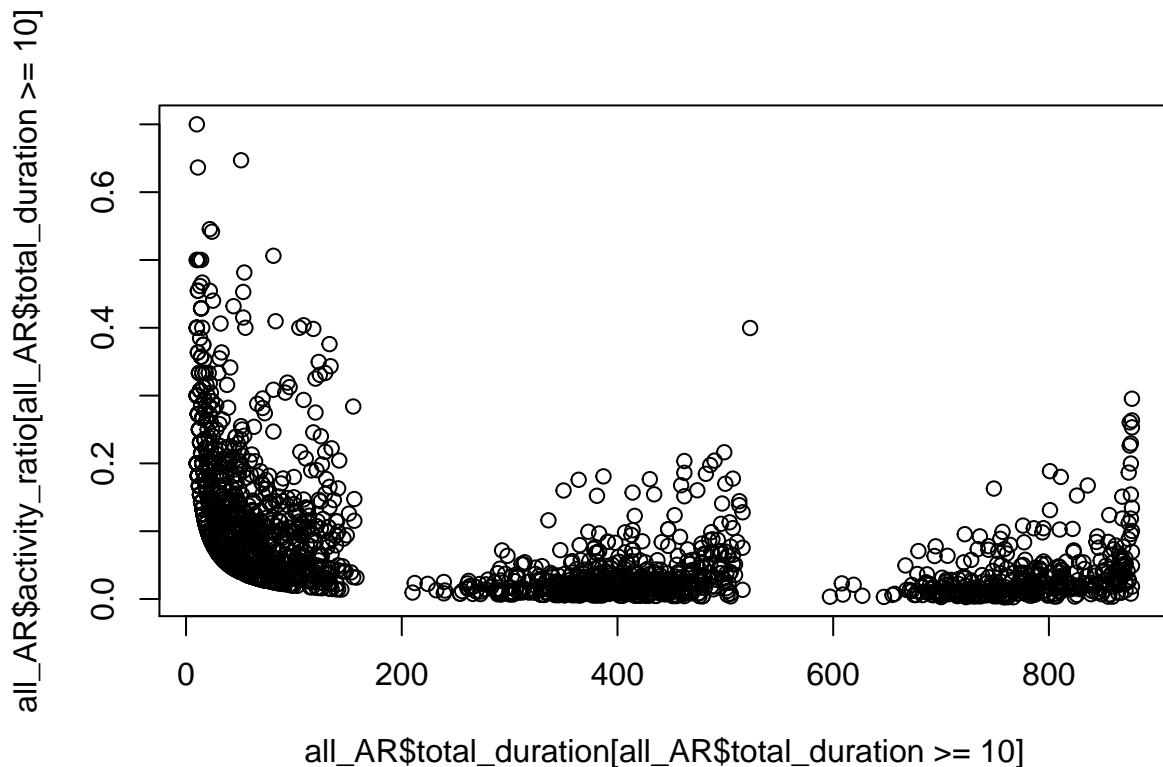


Both have nice distributions, though we can see the single record people in the ratio plot

```
# Plot activity_ratio against duration  
# Here we probably want to subset to avoid bias  
plot(all_AR$total_duration,  
      all_AR$activity_ratio)
```



```
plot(all_AR$total_duration[all_AR$total_duration >= 10],  
     all_AR$activity_ratio[all_AR$total_duration >= 10])
```



### Weekly devoted days

This is an adaptation of the *Daily Devoted Time* in (Ponciano and Brasileiro 2014) which is clearly not applicable to biological recording. Though Boakes *et al* 2016 don't attempt to use this measure I think the idea can be adapted by using days in a week (summer only) rather than hours in a day.

```
# Create a function
weeklyDevotedDays <- function(recorder_name,
                              data,
                              recorder_col = 'recorders',
                              date_col = 'date_start'){

  # check date column
  if(!inherits(data[, date_col], 'Date')){
    stop('Your date column is not a date')
  }

  # Get the recorders data
  data <- data[data[, recorder_col] == recorder_name, ]

  # Get unique dates as dates
  dates <- unique(data[, date_col])

  # Get all week_year combinations
  week_year <- paste(strftime(as.POSIXlt(dates), format = '%W'),
```

```

        format(dates, '%Y'), sep = '_')

# here are the counts
week_counts <- table(week_year)

# As these are counts taking the median is probably best
weekly_devoted_days <- median(week_counts)

return(data.frame(recorder = recorder_name,
                  median_weekly_devoted_days = weekly_devoted_days,
                  n_weeks = length(week_counts),
                  n_recs = sum(week_counts), row.names = NULL))
}

# Test on David and Tom
weeklyDevotedDays(data = summer_iRB, recorder_name = 'Roy, David')

```

```

##      recorder median_weekly_devoted_days n_weeks n_recs
## 1 Roy, David                2         60     131

```

```

weeklyDevotedDays(data = summer_iRB, recorder_name = 'August, Tom')

```

```

##      recorder median_weekly_devoted_days n_weeks n_recs
## 1 August, Tom                1         6       7

```

```

## David contributes more of his time than Tom ##

```

```

# Run for everyone
all_WDD <- do.call(rbind, lapply(X = unique(iRB$recorders),
                                FUN = weeklyDevotedDays,
                                data = summer_iRB))

```

```

# Lets have a look at some of these
head(all_WDD, 20)

```

```

##      recorder median_weekly_devoted_days n_weeks n_recs
## 1      Brookes , Anne                1.0     38     59
## 2      Burgoyne, Steve                2.0     18     47
## 3      Brown, Peter                  2.0     25     57
## 4      Rutherford, Joanna             1.0      7     11
## 5      Allan, David                   4.0     67    259
## 6      Millward, Martin               1.0      4      4
## 7      Foulkes-Arellano, Paul          1.0     13     18
## 8      Stewart, Tam                   3.0     65    231
## 9      Forbes, Andrew                 1.0      3      5
## 10     Richardson, Rosie              1.5      4      7
## 11     Partridge, Francesca           2.0     56    146
## 12     Card , Graeme                  1.5     12     22
## 13     Honey, Hawk                   1.0     39     72
## 14     Melzack, David                  1.5     12     30
## 15     Povall, Ed                     1.0      3      3
## 16     Goodwin, Paul                  2.0     26     65

```



## 17	Coulson, Joe	2.0	10	22
## 18	Bailey, Peggy	2.0	16	41
## 19	Roy, David	2.0	60	131
## 20	Woodley, Caroline	1.0	12	18

Clearly this metric is only really reliable when we have multiple weeks worth of data for an individual.

## Relative activity duration

This is a metric from Ponciano and Brasileiro 2014 which is also used in Boakes *et al* 2016 but I don't think can be applied to biological records since there is no official end date for a project: *"The ratio of days during which a volunteer I remains linked to the project in relation to the total number of days elapsed since the volunteer joined the project until the project is over"*

## Periodicity

There is a cluster of metrics that could be used to look at aspects of periodicity. The measure used in Ponciano and Brasileiro 2014 is 'variation in periodicity'; *"The standard deviation of the times elapsed between each pair of sequential active days"*. At the same time as calculating this I think there are another couple of metrics that might be of use. First, periodicity itself, i.e. *"The median time elapsed between each pair of sequential active days"*. Secondly, streak length, i.e. *"The average length of sequential active days"*

```
# Create a function to calculate the periodicity metrics
periodicity <- function(recorder_name,
                        data,
                        recorder_col = 'recorders',
                        date_col = 'date_start',
                        day_limit = 5){

  # check date column
  if(!inherits(data[, date_col], 'Date')){
    stop('Your date column is not a date')
  }

  # Get the recorders data
  data <- data[data[, recorder_col] == recorder_name, ]

  # Get unique dates as dates
  dates <- sort(unique(data[, date_col]))

  # we cannot calculate these metrics if people have very few
  # dates on which they record
  if(length(unique(dates)) < day_limit){

    # return
    return(data.frame(recorder = recorder_name,
                      periodicity = NA,
                      periodicity_variation = NA,
                      median_streak = NA,
                      sd_streak = NA,
                      max_streak = NA,
                      n_days = length(unique(dates))))
  }
}
```

```

} else {

  # Calculate the elapsed days between each date in sequence
  # this needs to be done within years
  elapses <- NULL

  for(year in unique(format(dates, '%Y'))){

    temp_dates <- dates[format(dates, '%Y') == year]

    # There must be at least 2 dates in a year
    if(length(temp_dates) > 1){
      temp_elapses <- sapply(1:(length(temp_dates)-1),
        FUN = function(x){
          return(as.numeric(temp_dates[x + 1] - temp_dates[x]))
        })

      elapses <- c(elapses, temp_elapses)

    }

  }

  # periodicity calculation
  periodicity <- median(elapses)

  # variation in periodicity
  periodicity_variation <- sd(elapses)

  # average streak length
  # Streaks are IDed by 1's
  non_streak <- length(elapses[elapses > 1])
  streaks <- rle(elapses)
  streaks_1 <- (streaks$lengths[streaks$value == 1]) + 1

  # Combine streaks and non-streaks
  streak_lengths <- c(rep(1, non_streak), streaks_1)

  # calculate ome metrics
  median_streak <- median(streak_lengths)
  sd_streak <- sd(streak_lengths)
  max_streak <- max(streak_lengths)

  # return
  return(data.frame(recorder = recorder_name,
    periodicity = periodicity,
    periodicity_variation = periodicity_variation,
    median_streak = median_streak,
    sd_streak = sd_streak,
    max_streak = max_streak,
    n_days = length(unique(dates))))

}

```

```

}

# Test on David and Tom
periodicity(data = summer_iRB, recorder_name = 'Roy, David')

##      recorder periodicity periodicity_variation median_streak sd_streak
## 1 Roy, David           2           3.196601           1 0.9699536
##   max_streak n_days
## 1           7    131

periodicity(data = summer_iRB, recorder_name = 'August, Tom')

##      recorder periodicity periodicity_variation median_streak sd_streak
## 1 August, Tom           30           17.81853           1 0.4472136
##   max_streak n_days
## 1           2      7

# David is a much more regular recorder than Tom with less
# variation in periodicity and a longer max streak though
# Tom has less days of data to work with

# Run for everyone
all_P <- do.call(rbind, lapply(X = unique(iRB$recorders),
                              FUN = periodicity,
                              data = iRB))

# Lets have a look at some of these
head(all_P, 20)[c(5,8,1),]

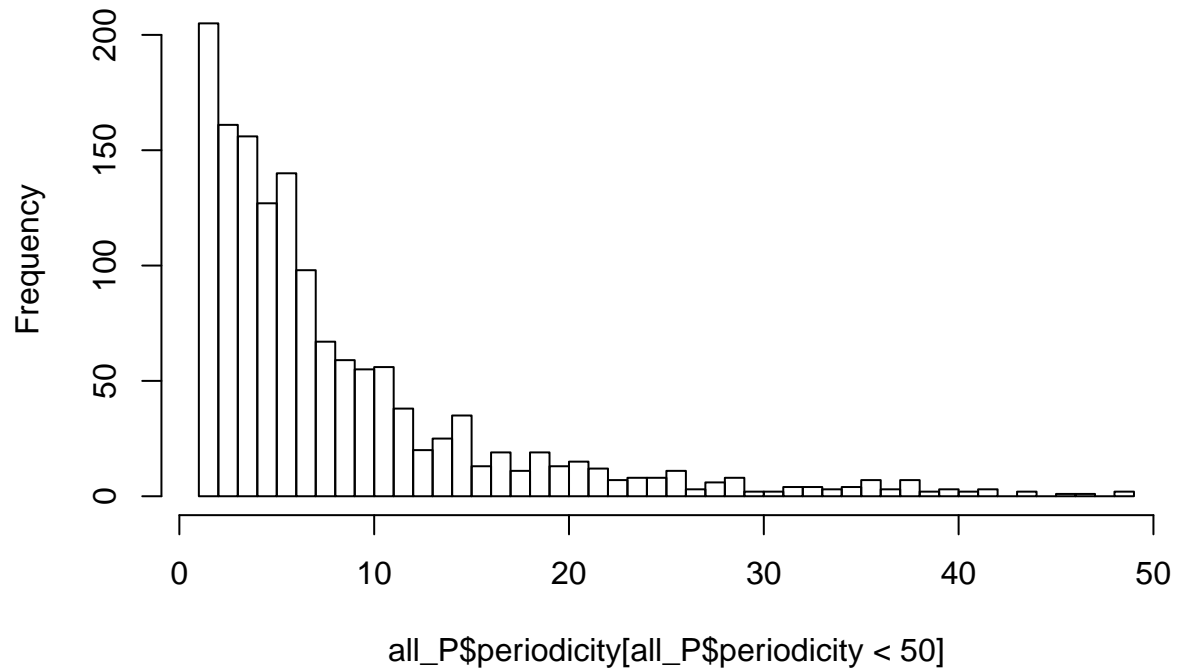
##      recorder periodicity periodicity_variation median_streak
## 5 Allan, David           1           2.693493           1
## 8 Stewart, Tam           1           2.237841           1
## 1 Brookes , Anne         5           8.101543           1
##   sd_streak max_streak n_days
## 5 2.0710394      16    378
## 8 1.8107359      15    272
## 1 0.5918027       4     70

# David a Tam are both very studious recorders with
# long max streaks and very low periodicity.
# Anne is less studious but still has a low periodicity

# Nice poisson dist. for periodicity
hist(all_P$periodicity[all_P$periodicity < 50],
     breaks = 50)

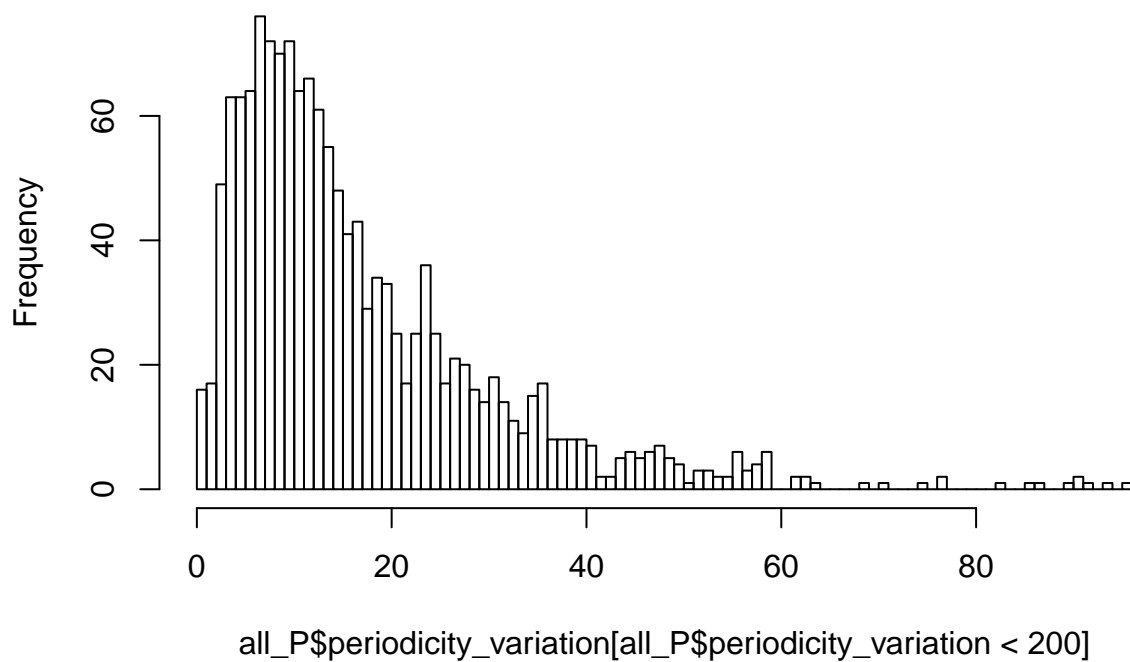
```

**Histogram of all\_P\$periodicity[all\_P\$periodicity < 50]**

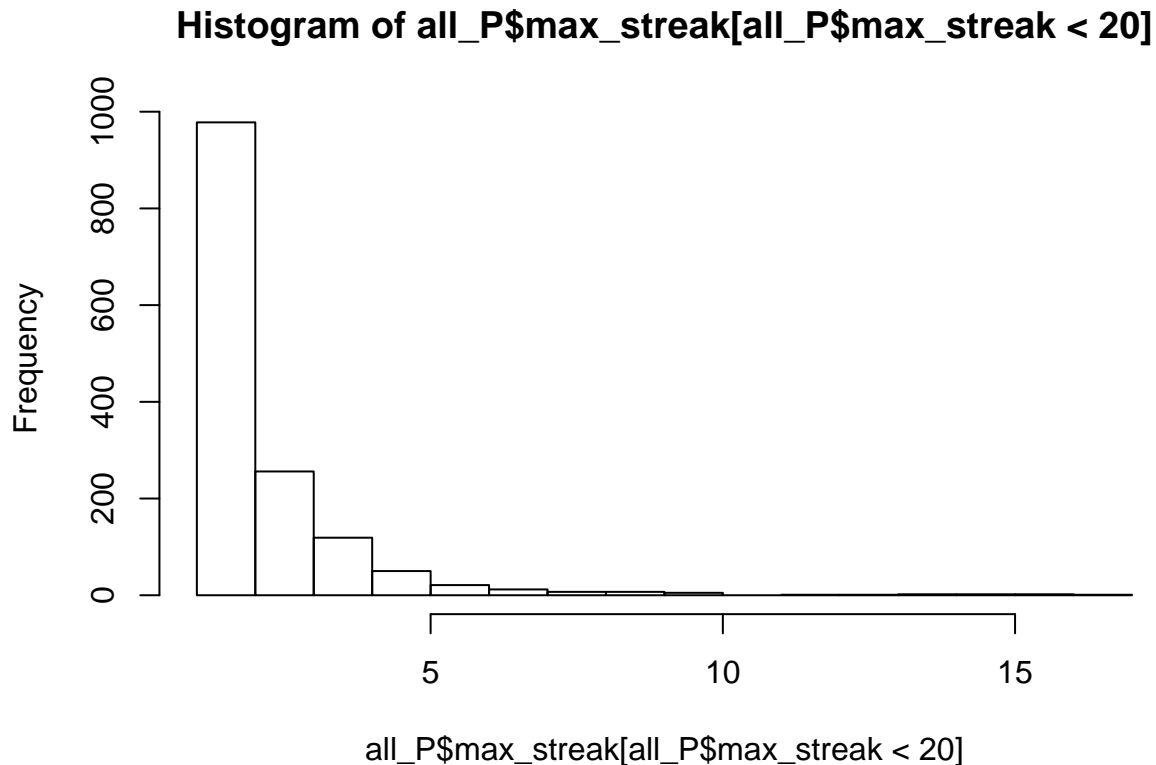


```
# Nice poisson dist. for periodicity_variation (long tail)  
hist(all_P$periodicity_variation[all_P$periodicity_variation < 200],  
      breaks = 100)
```

## Histogram of all\_P\$periodicity\_variation[all\_P\$periodicity\_variation < 200]



```
# Dist. of max_streak  
hist(all_P$max_streak[all_P$max_streak < 20],  
      breaks = 20)
```



By using the summer data only this analysis seems to be better than an earlier one that included all data. These metrics cannot be calculate for people who have only made one record. I have included a parameter `day_limit` to allow us to set a limit at which we calculate these metrics.

## Spatial Meterics

These metrics deal with the spatial distribution of records

### Area and heterogenity of recording

I think the first step for all of these metrics is to turn the points into a `SpatialPoints` object which will allow us to manipulate then more easily. Once we have done that we can calculate MCP (minimum convex polygons) around the points. We might want to change this method to a method that is less susceptible to outliers such as alpha hull (we can talk to Colin about this). Here I use 95% MCP as the total recording area (hopefully removing outliers), and use the ratio of 95%:50% as a measure of heterogeneity.

```
# Function takes data and username and returns spatial metrics
spatial_behaviour <- function(data, recorder_name,
                              latitude_col, longitude_col,
                              recorder_col = 'recorders',
                              upper_percentile = 95,
                              lower_percentile = 50){

  if(is.factor(recorder_name)){
    recorder_name <- as.character(recorder_name)
  }
}
```

```

}

n_row <- nrow(iRB[iRB[,recorder_col] == recorder_name, ])

if(n_row >= 5){

  # Convert to SpatialPoints
  spPoints_LL <- SpatialPoints(iRB[iRB[,recorder_col] == recorder_name,
                                   c(longitude_col, latitude_col)])

  # Data is lat long
  proj4string(spPoints_LL) <- CRS("+init=epsg:4326")

  # Convert to Eastings Northings to get meters on X and Y
  spPoint_UK <- spTransform(spPoints_LL, "+init=epsg:27700")

  # Calculate the larger MCP
  mcp_poly_upper <- mcp(spPoint_UK,
                        percent = upper_percentile,
                        unin = 'm',
                        unout = 'km2')

  # Calculate the smaller MCP
  mcp_poly_lower <- mcp(spPoint_UK,
                        percent = lower_percentile,
                        unin = 'm',
                        unout = 'km2')

  return(list(recorder = recorder_name,
              spPoint_UK = spPoint_UK,
              mcp_poly_upper = mcp_poly_upper,
              mcp_poly_lower = mcp_poly_lower,
              upper_area = mcp_poly_upper$area,
              lower_area = mcp_poly_lower$area,
              ratio = mcp_poly_lower$area/mcp_poly_upper$area,
              n = n_row))
} else {
  return(list(recorder = recorder_name,
              spPoint_UK = NA,
              mcp_poly_upper = NA,
              mcp_poly_lower = NA,
              upper_area = NA,
              lower_area = NA,
              ratio = NA,
              n = n_row))
}

}

# Test on one recorder
David_spatial <- spatial_behaviour(data = iRB, recorder_name = 'Roy, David',
                                   latitude_col = 'lat', longitude_col = 'st_x')

# Function for plotting records

```

```

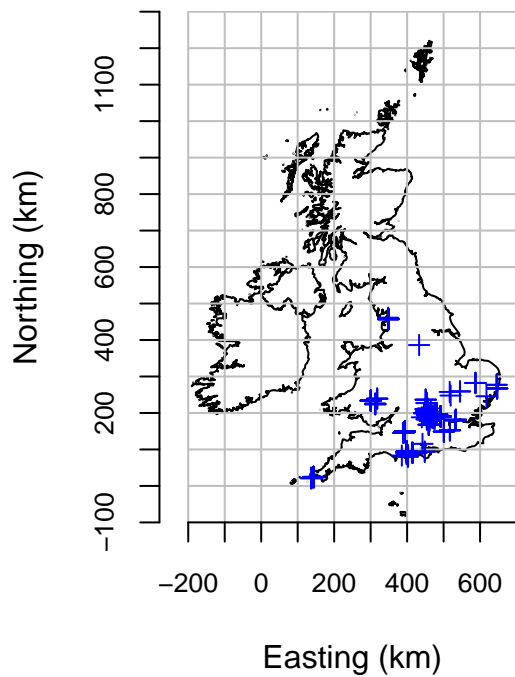
plot_ratio <- function(data){
  par(mfrow = c(1,2))
  data(UK)
  plot_GIS(UK, new.window = FALSE, main = 'Distribution of records')
  points(data$spPoint_UK, pch = 3, col = 'blue')

  # Plot David's heat map
  plot(data$spPoint_UK,
        main = paste(data$recorder, '-', 'Ratio:', round(data$ratio, 4)),
        col = 'blue')
  upper_polygon <- data$mcp_poly_upper@polygons[[1]]@Polygons[[1]]@coords
  polygon(x = upper_polygon[,1],
          y = upper_polygon[,2])
  lower_polygon <- data$mcp_poly_lower@polygons[[1]]@Polygons[[1]]@coords
  polygon(x = lower_polygon[,1],
          y = lower_polygon[,2],
          col = 'red', border = 'red')
  par(mfrow = c(1,1))
}

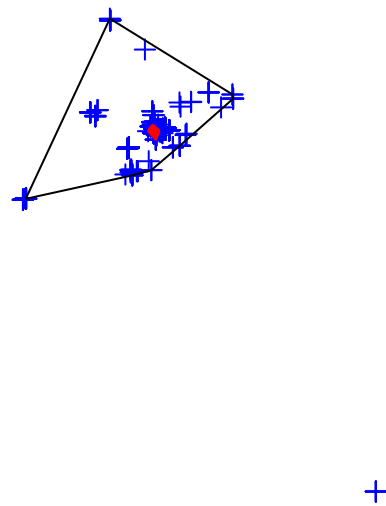
# Plot
plot_ratio(data = David_spatial)

```

**Distribution of records**



**Roy, David – Ratio: 0.0051**



## NOTE DAVID HAS A RECORD FROM OUTSIDE THE UK ##



```

# Apply to all recorders
all_spatial <- lapply(unique(iRB$recorders), FUN = function(x){
  recorder_info <- spatial_behaviour(data = iRB, recorder_name = x,
                                     latitude_col = 'lat', longitude_col = 'st_x')
  return(data.frame(recorder = recorder_info$recorder,
                    upper_area = recorder_info$upper_area,
                    lower_area = recorder_info$lower_area,
                    ratio = recorder_info$ratio,
                    n = recorder_info$n))
})

# combine results
temp <- do.call(rbind, all_spatial)
temp <- temp[temp$n > 400, ]

# Lets have a look at some people who have recorded a lot
temp[order(temp$ratio, decreasing = TRUE),]

```

	recorder	upper_area	lower_area	ratio	n
## 11	Partridge, Francesca	5.176381e+03	2.414166e+03	0.4663809300	1418
## 52	Cornish, Stephen	3.534308e+00	9.106945e-01	0.2576726378	487
## 180	Limb, Ken	3.042189e+04	7.577886e+03	0.2490932324	622
## 395	Atkin, Paul	1.393205e+03	3.223848e+02	0.2313978875	615
## 139	Hunter, Amands	7.531823e+02	1.246409e+02	0.1654856950	1090
## 104	Leaver, Kim	1.394622e+03	2.193449e+02	0.1572790538	537
## 26	fenn, paul	5.583057e+03	8.487535e+02	0.1520230771	2503
## 65	Gillie, Tony	1.750010e+03	2.097561e+02	0.1198599351	1112
## 339	Bowles, Nick	4.155985e+03	3.545317e+02	0.0853062848	590
## 256	Cowton, Keith	2.109076e+04	1.119724e+03	0.0530907308	445
## 113	Hill, Brian	7.170905e+03	3.793400e+02	0.0528998784	851
## 5	Allan, David	1.471503e+04	6.918175e+02	0.0470143525	3180
## 39	Warren, Martin	3.863468e+04	1.337492e+03	0.0346189363	2434
## 72	Jones, Dave	2.767527e+01	9.346723e-01	0.0337728352	2207
## 109	Shanks, Scott	2.523051e+04	8.281931e+02	0.0328250625	513
## 103	Pennington, Robert	6.234135e+03	1.838310e+02	0.0294878026	969
## 1356	Saville, Simon	2.969962e+04	8.676054e+02	0.0292126767	441
## 383	Steele, Andrew	9.131555e+04	2.632030e+03	0.0288234530	563
## 123	Cox, Steve	4.447586e+04	1.265539e+03	0.0284545070	991
## 8	Stewart, Tam	2.886784e+04	8.000475e+02	0.0277141435	1811
## 175	Sims, Clive	2.359346e+04	6.345611e+02	0.0268956338	864
## 41	Lonsdale, Liz and Steve	1.536766e+05	3.975898e+03	0.0258718467	542
## 523	Shersby, Megan	3.790063e+04	9.782456e+02	0.0258108020	478
## 43	Newbould, John	7.404497e+04	1.879715e+03	0.0253861332	1001
## 488	Kilbey, Dave	3.198760e+04	6.217184e+02	0.0194362359	780
## 45	Sell, Claire	7.754820e+02	1.501250e+01	0.0193589276	555
## 197	Lunnon, Marie	6.657662e+01	1.256184e+00	0.0188682491	444
## 96	Checkley, Graham	1.240849e+03	2.166370e+01	0.0174587706	1813
## 143	Fox, Richard	3.168087e+04	4.871148e+02	0.0153756723	1147
## 19	Roy, David	1.065308e+05	5.448197e+02	0.0051142000	615
## 78	shilland, ewan	1.519489e+05	6.898383e+02	0.0045399371	1636
## 87	Dawson, Steve	1.135666e+03	4.548707e+00	0.0040053224	789
## 140	Austin, David	4.573147e+03	1.795774e+01	0.0039267802	441
## 100	Ford, Rachel	7.010182e+01	9.404533e-02	0.0013415532	431

```
## 158          Harley, Ross 1.873400e+05 7.630491e+01 0.0004073071 682
```

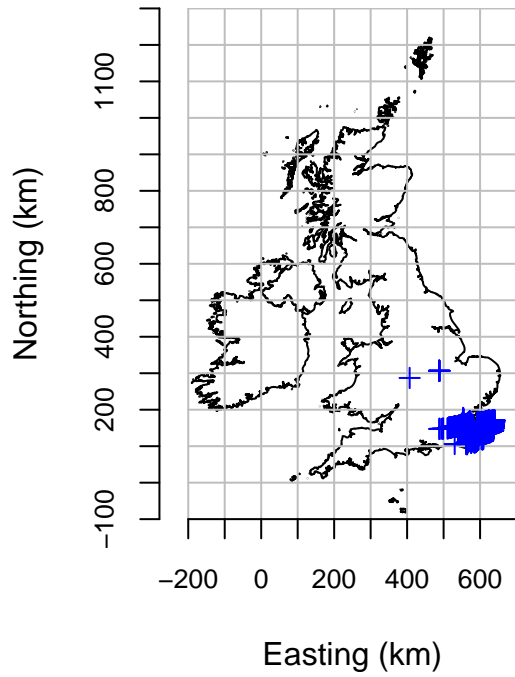
Lets have a look at two people with very different ratios

```
# Get the names of top and bottom
temp <- temp[order(temp$ratio, decreasing = TRUE),]

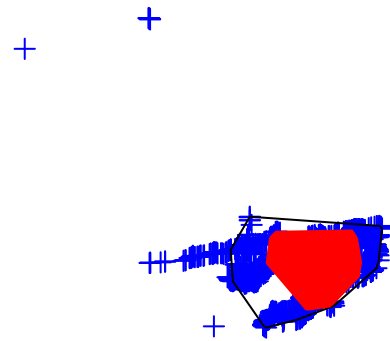
top <- as.character(head(temp$recorder, 1))
bottom <- as.character(tail(temp$recorder, 1))

# Plot the top and bottom ratio recorder
for(i in c(top, bottom)){
  top_d <- spatial_behaviour(data = iRB,
                             recorder_name = i,
                             latitude_col = 'lat',
                             longitude_col = 'st_x')
  plot_ratio(data = top_d)
}
```

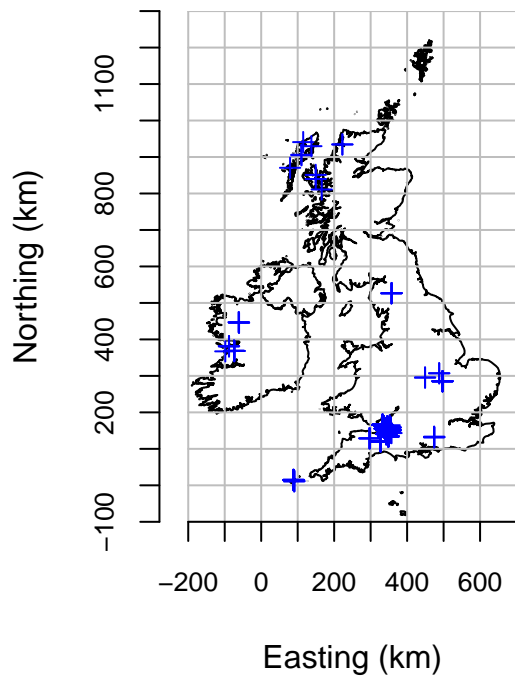
**Distribution of records**



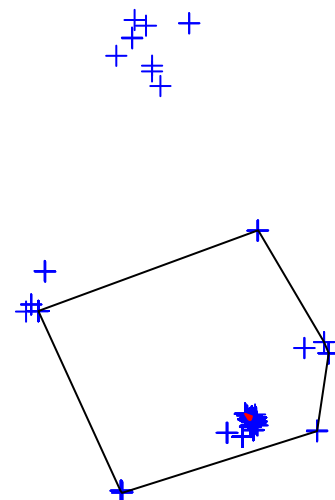
**Partridge, Francesca – Ratio: 0.46**



**Distribution of records**



**Harley, Ross – Ratio: 4e-04**



## Taxonomic Metrics

These metric relate the the species that people record

### Taxonomic Breadth

This is simply a measure of the proportion of taxa a person has recorded. Note this is going to be correlated to the number of records.

```
taxa_breadth <- function(data, recorder_name,
                          sp_col = 'preferred_taxon',
                          recorder_col = 'recorders'){

  data_rec <- data[data[,recorder_col] == recorder_name, c(sp_col, recorder_col)]

  return(data.frame(recorder = recorder_name,
                    taxa_breadth = length(unique(data_rec[,sp_col])),
                    taxa_prop = length(unique(data_rec[,sp_col]))/length(unique(data[,sp_col])),
                    n = nrow(data_rec)))
}

taxa_breadth <- do.call(rbind, lapply(unique(iRB$recorders), FUN = taxa_breadth, data = iRB))

temp <- taxa_breadth[taxa_breadth$n > 400, ]

# Lets have a look at some people who have recorded a lot
temp[order(temp$taxa_prop, decreasing = TRUE),]
```

##	recorder	taxa_breadth	taxa_prop	n
## 39	Warren, Martin	52	0.6265060	2434
## 5	Allan, David	51	0.6144578	3180
## 103	Pennington, Robert	49	0.5903614	969
## 113	Hill, Brian	48	0.5783133	851
## 1356	Saville, Simon	48	0.5783133	441
## 123	Cox, Steve	47	0.5662651	991
## 175	Sims, Clive	47	0.5662651	864
## 143	Fox, Richard	46	0.5542169	1147
## 158	Harley, Ross	45	0.5421687	682
## 383	Steele, Andrew	45	0.5421687	563
## 256	Cowton, Keith	42	0.5060241	445
## 395	Atkin, Paul	42	0.5060241	615
## 26	fenn, paul	41	0.4939759	2503
## 180	Limb, Ken	41	0.4939759	622
## 488	Kilbey, Dave	41	0.4939759	780
## 87	Dawson, Steve	40	0.4819277	789
## 65	Gillie, Tony	39	0.4698795	1112
## 523	Shersby, Megan	38	0.4578313	478
## 19	Roy, David	37	0.4457831	615
## 41	Lonsdale, Liz and Steve	37	0.4457831	542
## 78	shilland, ewan	36	0.4337349	1636
## 339	Bowles, Nick	36	0.4337349	590
## 43	Newbould, John	33	0.3975904	1001
## 11	Partridge, Francesca	32	0.3855422	1418

## 45	Sell, Claire	32	0.3855422	555
## 139	Hunter, Amands	31	0.3734940	1090
## 8	Stewart, Tam	29	0.3493976	1811
## 197	Lunnon, Marie	28	0.3373494	444
## 109	Shanks, Scott	26	0.3132530	513
## 104	Leaver, Kim	24	0.2891566	537
## 72	Jones, Dave	23	0.2771084	2207
## 96	Checkley, Graham	22	0.2650602	1813
## 140	Austin, David	22	0.2650602	441
## 52	Cornish, Stephen	19	0.2289157	487
## 100	Ford, Rachel	15	0.1807229	431

## Species Rarity

We want to capture the rarity of the species that people record. For example are they just recording the common species or are they only recording the rare ones, or perhaps they are recording everything. Since we don't know the real frequency distribution we can only compare people to the global average in the dataset. We can look to see what the distribution of species rank for each recorder is and how this compares to all records. A recorder only interested in rare species will have a median rank higher than the average. A recorder only recording common species will have a value lower than the average.

```
# Lets look at a recorder
species_rank <- function(data, recorder_name,
                          sp_col = 'preferred_taxon',
                          recorder_col = 'recorders'){

  data <- data[,c(sp_col, recorder_col)]
  rank_species <- rank(abs(table(data[,sp_col]))-max(table(data[,sp_col]))))
  sp_counts <- table(data[,sp_col])

  rank_reps <- rep(rank_species, sp_counts)
  grand_median <- median(rank_reps)
  grand_sd <- sd(rank_reps)

  recorder_data <- data[data[,recorder_col] == recorder_name,]
  recorder_data$rank <- rank_species[recorder_data[,sp_col]]

  return(data.frame(recorder = as.character(recorder_name),
                    median = median(recorder_data$rank),
                    median_diff = median(recorder_data$rank) - grand_median,
                    stdev = sd(recorder_data$rank),
                    n = nrow(recorder_data)))
}

rarity_preference <- do.call(rbind,
                             lapply(unique(iRB$recorders),
                                     FUN = species_rank,
                                     data = iRB))

temp <- rarity_preference[rarity_preference$n > 400, ]

# Lets have a look at some people who have recorded a lot
temp[order(temp$median_diff, decreasing = TRUE),]
```

##	recorder	median	median_diff	stdev	n
## 1356	Saville, Simon	13	5	12.191833	441
## 256	Cowton, Keith	12	4	10.283900	445
## 39	Warren, Martin	11	3	10.754206	2434
## 175	Sims, Clive	11	3	10.132960	864
## 339	Bowles, Nick	10	2	8.557264	590
## 395	Atkin, Paul	10	2	9.738285	615
## 523	Shersby, Megan	10	2	8.613459	478
## 8	Stewart, Tam	9	1	10.764394	1811
## 19	Roy, David	9	1	9.647095	615
## 26	fenn, paul	9	1	8.779256	2503
## 43	Newbould, John	9	1	8.245020	1001
## 45	Sell, Claire	9	1	8.912894	555
## 65	Gillie, Tony	9	1	8.645367	1112
## 103	Pennington, Robert	9	1	9.100094	969
## 109	Shanks, Scott	9	1	9.482688	513
## 113	Hill, Brian	9	1	10.226885	851
## 139	Hunter, Amands	9	1	7.199181	1090
## 158	Harley, Ross	9	1	9.410956	682
## 180	Limb, Ken	9	1	9.165788	622
## 197	Lunnon, Marie	9	1	7.004225	444
## 41	Lonsdale, Liz and Steve	8	0	8.646054	542
## 78	shilland, ewan	8	0	8.303214	1636
## 96	Checkley, Graham	8	0	6.931797	1813
## 104	Leaver, Kim	8	0	6.082150	537
## 143	Fox, Richard	8	0	9.681677	1147
## 383	Steele, Andrew	8	0	9.108308	563
## 488	Kilbey, Dave	8	0	9.170174	780
## 87	Dawson, Steve	7	-1	7.926813	789
## 100	Ford, Rachel	7	-1	5.281118	431
## 123	Cox, Steve	7	-1	9.048282	991
## 5	Allan, David	6	-2	8.643921	3180
## 11	Partridge, Francesca	6	-2	6.888191	1418
## 72	Jones, Dave	6	-2	4.862982	2207
## 52	Cornish, Stephen	5	-3	5.081520	487
## 140	Austin, David	5	-3	5.474312	441

Here `median_diff` gives the difference between the grand median for all records and the recorders median. This suggests **Saville, Simon** prefers to record rare species and **Cornish, Stephen** prefers to record common species.

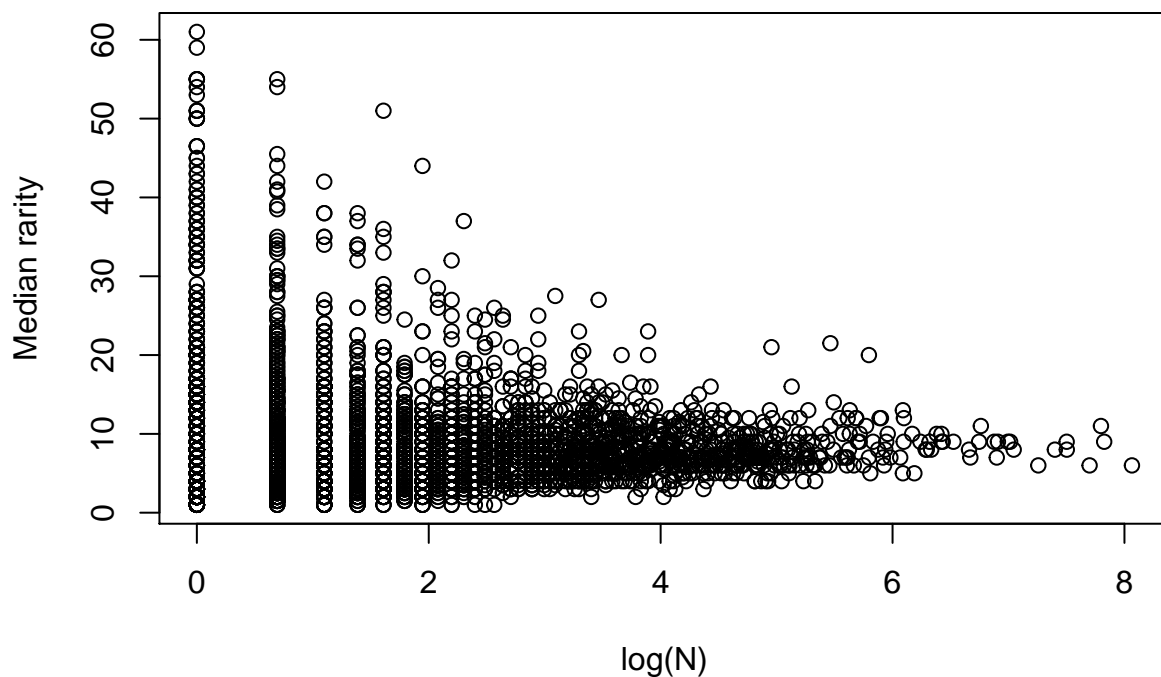
This could be correlated to the number of records.

```
mod <- glm(median ~ log(n), data = rarity_preference, family = 'quasipoisson')
summary(mod)
```

```
##
## Call:
## glm(formula = median ~ log(n), family = "quasipoisson", data = rarity_preference)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -3.7397  -1.6376  -0.4224   0.7604  10.7394
##
```

```
## Coefficients:
##             Estimate Std. Error t value Pr(>|t|)
## (Intercept)  2.334807   0.018526 126.030  <2e-16 ***
## log(n)       -0.070761   0.008457  -8.367  <2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for quasipoisson family taken to be 5.26387)
##
## Null deviance: 17611  on 3944  degrees of freedom
## Residual deviance: 17232  on 3943  degrees of freedom
## AIC: NA
##
## Number of Fisher Scoring iterations: 5
```

```
plot(log(rarity_preference$n),
     rarity_preference$median,
     xlab = 'log(N)',
     ylab = 'Median rarity')
```



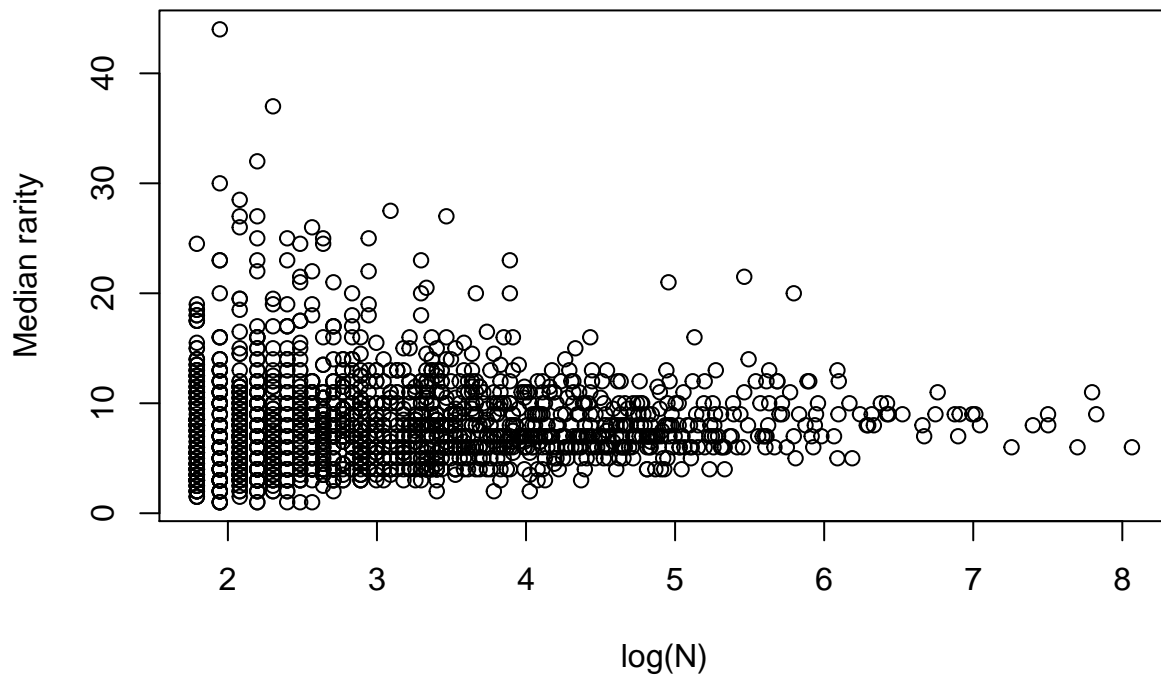
There is a significant negative relationship. The more records you make the lower your median value. This could be a result of the fact that people who make only a few records record rare stuff?

```
rarity_preference_above <- rarity_preference[rarity_preference$n > 5, ]
mod <- glm(median ~ log(n), data = rarity_preference_above, family = 'quasipoisson')
summary(mod)
```

```
##
## Call:
## glm(formula = median ~ log(n), family = "quasipoisson", data = rarity_preference_above)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -3.1729  -0.9646  -0.2134   0.6260   8.7863
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  2.076269   0.034427  60.310  <2e-16 ***
## log(n)       0.007479   0.010513   0.711   0.477
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for quasipoisson family taken to be 2.087498)
##
##      Null deviance: 3474.5  on 1877  degrees of freedom
## Residual deviance: 3473.5  on 1876  degrees of freedom
## AIC: NA
##
## Number of Fisher Scoring iterations: 4
```

```
plot(log(rarity_preference_above$n),
     rarity_preference_above$median,
     xlab = 'log(N)',
     ylab = 'Median rarity')
```





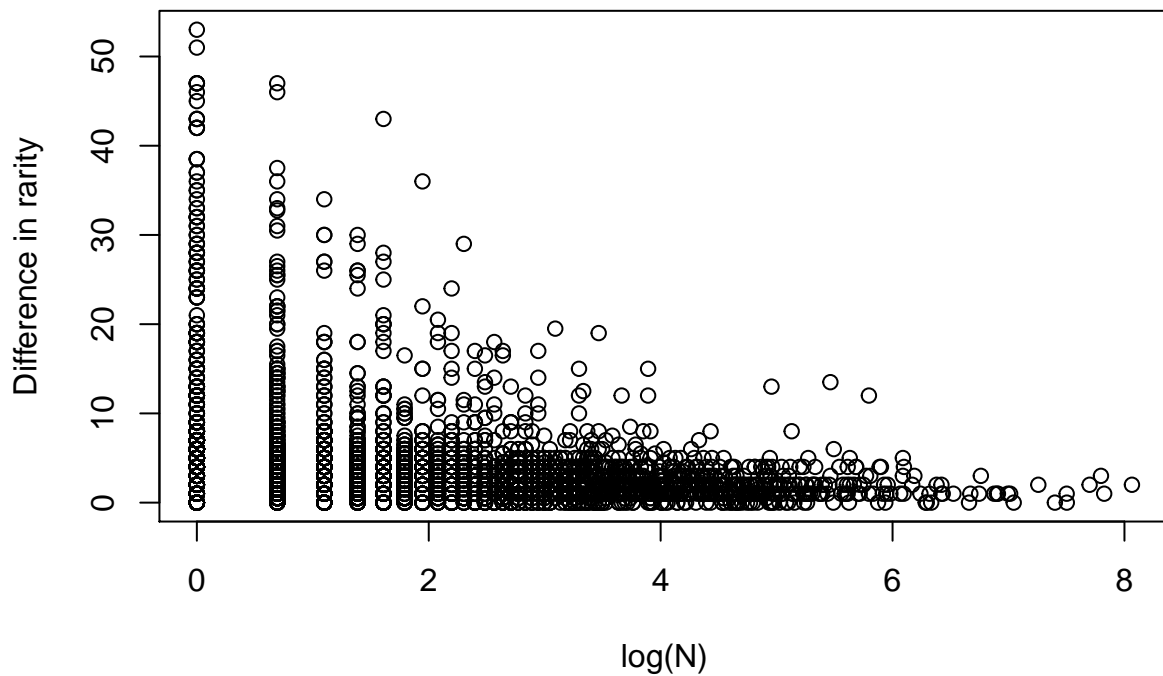
Okay, the relationship falls down once we get rid of the people who only record a few species. I suggest this metric not be estimates for people who contribute only a few records. The relationship might actually be between deviation from the median and  $n$ .

```
rarity_preference$median_diff_abs <- abs(rarity_preference$median_diff)
mod <- glm(median_diff_abs ~ log(n), data = rarity_preference, family = 'quasipoisson')
summary(mod)
```

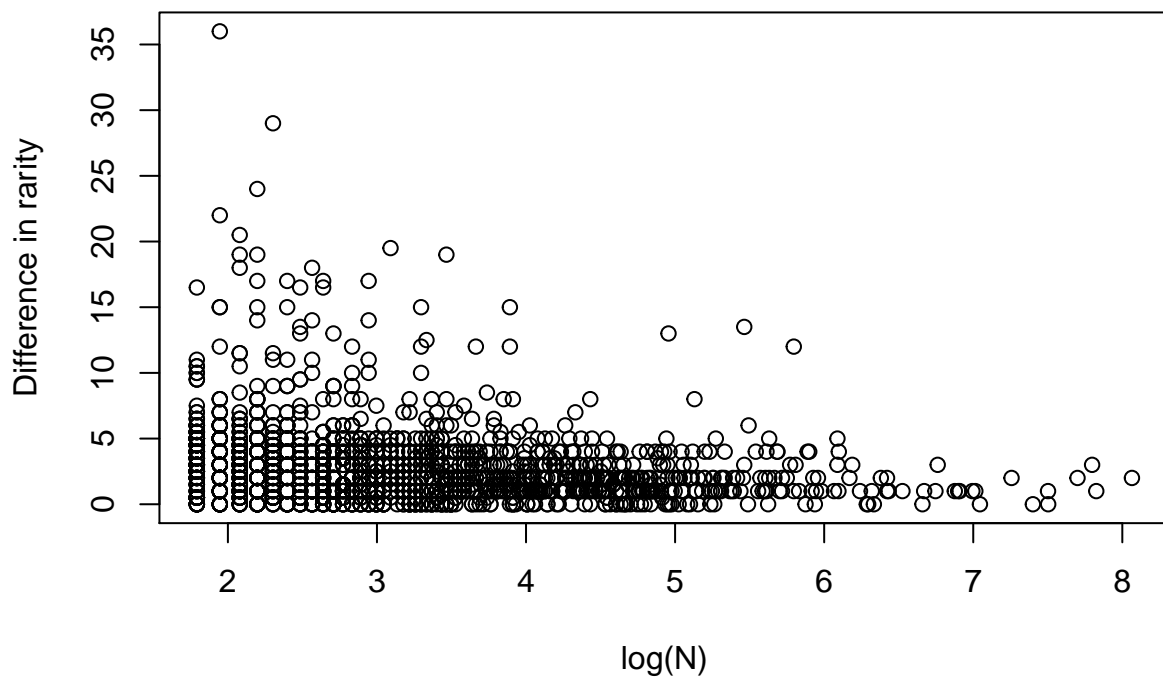
```
##
## Call:
## glm(formula = median_diff_abs ~ log(n), family = "quasipoisson",
##      data = rarity_preference)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -3.8262  -1.3717  -0.5038   0.3857  10.8928
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  1.99062    0.02414   82.47  <2e-16 ***
## log(n)       -0.31472    0.01388  -22.68  <2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for quasipoisson family taken to be 5.259472)
##
```

```
## Null deviance: 18546 on 3944 degrees of freedom
## Residual deviance: 15454 on 3943 degrees of freedom
## AIC: NA
##
## Number of Fisher Scoring iterations: 5
```

```
plot(log(rarity_preference$n),
     rarity_preference$median_diff_abs,
     xlab = 'log(N)',
     ylab = 'Difference in rarity')
```



```
plot(log(rarity_preference$n[rarity_preference$n > 5]),
     rarity_preference$median_diff_abs[rarity_preference$n > 5],
     xlab = 'log(N)',
     ylab = 'Difference in rarity')
```



The more records you record the less you deviate from the median. This is probably because you only get extreme values where the sample size is small.