Explaining the rc Functions

Mark White | markhwhiteii@gmail.com

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I will discuss the rc ("Records Count") functions for feeds (feedrc) and facilities (facilityrc) in this document. The functions can be loaded by running the rc.R document, and the rcGuide.pdf file explains how to operate it. In case the CDC changes variable names, connections change, people are interested in doing something different with the code, etc., I will explain here line-by-line what the code means. I will first present a segment of code and then explain it below.

The feedro function

```
feedrc <- function(channel, visit, arrived, month, year) {</pre>
```

This first line names the function feedrc and specifies the five different arguments that it will take (see rcGuide.pdf for a description of these arguments).

```
suppressPackageStartupMessages(require(lubridate))
suppressPackageStartupMessages(require(data.table))
suppressPackageStartupMessages(require(dplyr))
suppressPackageStartupMessages(require(Hmisc))
```

These lines load the required packages if they are not already loaded. Some packages give a bunch of annoying messages when they are loaded, so I suppressed these for this function.

```
if (visit==TRUE) {
```

This code chunk will only be evaluated if the user specifies they want to get records for patient visits during that month. If not, it skips to the next if statement.

This code chunk generates a SQL query and pastes in the month and year that the user has specified, using the RODBC connection specified with the channel argument. This data is placed into a data frame called dv (for "data visits").

```
dv$Visit_Date <- gsub(" UTC", "",
floor_date(ymd_hms(dv$C_Visit_Date_Time), unit="day"))</pre>
```

When we pull the date down from the database, it includes hours, minutes, and seconds. We want to generalize this to the day, and this is what this line does. It takes the variable—dv\$C_Visit_Date_Time—and tells R that it is in the format of ymd_hms (year, month, day, hours, minutes, seconds). I use the floor_date() function to round all times down to the

nearest unit="day". This function will automatically add a time zone to the end. I do not want this time zone information, so I simply delete it using the gsub() function. This function says I want to take the pattern "UTC" and replace it with nothing (i.e, "").

```
outv <- setDT(as.data.frame.matrix(table(dv$Visit_Date, dv$Feed_Name)),
keep.rownames = TRUE)
}</pre>
```

Here, I take a frequency table of visit dates (dv\$Visit_Date) and feeds (dv\$Feed_Name), I save it as a data frame matrix, and then I set it as a data table, with the row names (dates) now stored as the first variable. It saves this into a new data frame, called outv (which stands for "output visits"). The last curly bracket means that I have closed the if statement. If the user did not specify visit as TRUE, then it has skipped over this whole thing.

This code chunk runs if the user specified TRUE for wanting the records that arrived during the month. This is the same exact code as above, except the data and output are suffixed with "a" (which stands for "arrived"), and we are now pulling down and working with Arrived_Data_Time instead of C_Visit_Date_Time.

```
if (visit==TRUE & arrived==FALSE) {
    setnames(outv, 1, "Visit_Date")
    return(outv)
```

This code chunk is run if the user asked for visits *only*. The first column is named "Visit_Date", and the output is returned to the user.

```
} else if (visit==FALSE & arrived==TRUE) {
    setnames(outa, 1, "Arrived_Date")
    return(outa)
```

This chunk does the same as the previous chunk of code, except it runs if the user asked for arrived *only*.

```
} else if (visit==TRUE & arrived==TRUE) {
    setnames(outa, 1, "Date")
    setnames(outv, 1, "Date")
    outav <- full_join(outa, outv, by="Date", suffix=c("_A", "_V"))
    outav[is.na(outav)] <- 0
    return(outav[order(outav$Date),c(colnames(outav)[1],
sort(colnames(outav)[2:length(colnames(outav))]))])</pre>
```

This runs if the user asked for both visits and arrivals. It renames the first column to "Date", and then the data are joined (using full_join()). The suffix argument is where the "_A" and "_V" are attached to the end of the column names. The next line specifies any missing values as a zero, and the last line returns the output. The arguments inside the brackets say that we want the rows ordered by Date, and the columns ordered with the data first and the rest done alphabetically.

```
} else {
    stop(cat("Please specify visit and/or arrived as TRUE."))
}
```

This last piece of code runs if the user specified FALSE for both visit and arrived.

The facilityre function

Most of the code here is the same as above, so I will only go into detail where it changes.

All of this code is the same as above, except that we are now pulling down C_Biosense_Facility_ID instead of Feed_Name.

The Processed table does not have the facility name, but the MFT table does. This code joins the data with another data frame that has two columns: the facility ID and name from the MFT table. It is a left_join, so it drops all rows from the MFT data frame that is not in the Processed data we first pulled down (i.e., it does not include rows for facilities that were not represented in the original data set we pulled down).

```
dv$Facility_Name <- droplevels(dv$Facility_Name)</pre>
```

Since Facility_Name was saved as a factor, it wants to return values for *all* of the facilities, even if they aren't found in the data set. We simply drop unused levels here, so that we don't have a bunch of zeros in our data frame from facilities that aren't in production or didn't report any records during the time frame we are looking at.

```
namekey <- unique(dv[,c("C Biosense Facility ID","Facility Name")])</pre>
```

This line generates a key for IDs and names: It takes the unique parts of the list of IDs matched with names (i.e., the facility ID and name are only represented once). We will use this in a second to match IDs with facility names

```
outv <- setDT(as.data.frame.matrix(table(dv$Visit_Date,
dv$C_Biosense_Facility_ID)), keep.rownames = TRUE)</pre>
```

Just like in the feedrc function, this creates the output we want.

```
for (i in 2:length(colnames(outv))) {
    id <- colnames(outv)[i]
    label(outv[[id]]) <-
as.character(namekey[which(namekey$C_Biosense_Facility_ID==id), "Facility_Name"])
    }
}</pre>
```

This is a for loop that performs an action for every column *except* for the first one (which is always date). It takes the ID number from the name of columns and saves it into an object called id. Then the label for this column is set to the Facility_Name that is on the same row as the ID number in the namekey. The curly brackets end the for loop and then end the if statement, respectively.

```
if (arrived==TRUE) {
    da <- sqlQuery(channel, paste0("SELECT Arrived_Date_Time,</pre>
C Biosense Facility ID
                                     FROM KS PR PRocessed
                                    WHERE MONTH(Arrived Date Time) = ", month,
" AND YEAR(Arrived_Date_Time) = ", year))
    da$Arrived_Date <- gsub(" UTC", "",</pre>
floor_date(ymd_hms(da$Arrived_Date_Time), unit="day"))
    outa <- setDT(as.data.frame.matrix(table(da$Arrived Date,
da$C Biosense Facility ID)), keep.rownames = TRUE)
    da <- left join(da, sqlQuery(channel, "SELECT C Biosense Facility ID,
Facility Name FROM KS MFT"),
                     by="C Biosense Facility ID")
    da$Facility_Name <- droplevels(da$Facility_Name)</pre>
    namekey <- unique(da[,c("C Biosense Facility ID", "Facility Name")])</pre>
    outa <- setDT(as.data.frame.matrix(table(da$Arrived_Date,</pre>
da$C Biosense Facility ID)), keep.rownames = TRUE)
    for (i in 2:length(colnames(outa))) {
      id <- colnames(outa)[i]</pre>
```

```
label(outa[[id]]) <-
as.character(namekey[which(namekey$C_Biosense_Facility_ID==id), "Facility_Name
"])
     }
}</pre>
```

This code chunk is the same as above, except it runs for the Arrived_Date_Time, and only will be executed if the user specifies TRUE for arrived.

```
if (visit==TRUE & arrived==FALSE) {
    setnames(outv, 1, "Visit_Date")
    return(outv)
  } else if (visit==FALSE & arrived==TRUE) {
    setnames(outa, 1, "Arrived_Date")
    return(outa)
  } else if (visit==TRUE & arrived==TRUE) {
    setnames(outa, 1, "Date")
    setnames(outv, 1, "Date")
    outav <- full_join(outa, outv, by="Date", suffix=c("_A", "_V"))</pre>
    outav[is.na(outav)] <- 0</pre>
    return(outav[order(outav$Date),c(colnames(outav)[1],
sort(colnames(outav)[2:length(colnames(outav))]))])
  } else {
    stop(cat("Please specify visit and/or arrived as TRUE."))
  }
}
```

Lastly, all of this code is the same as it was for the feedrc function; it is simply preparing the output and returning it to the user.