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fix.with <- function(xx = febl6WithAll.df, out.xls = "MissingWith.xls"){
### Purpose:- Slightly more generic
### -----
### Modified from:- fix.septLBAMwith
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### Arguments:-
### -----
### Author:- Patrick Connolly, Date:- 17 Feb 2016, 16:02
### -----
### Revisions:-

require(dplyr)

if(class(xx$Date) != "Date")
  xx <- within(xx, Date <- as.Date(as.character(Date), format = "%d/%m/%Y"))
## xx <- within(xx, Test <- as.numeric((Efnom)))
## xx <- within(xx, UC <- unclass(Efnom))
xx <- within(xx, Efnom <- as.numeric(as.character(Efnom)))
## xx <- xx[!is.na(xx$Efnom),]
xx <- within(xx, HC <- is.na(Efnom)) # no CO2 (handling control)
xx <- within(xx, Efnom[is.na(Efnom)] <- 0) # no EF either
xx <- within(xx, Dead[is.na(Dead)] <- 0) # one empty cell should be zero
is.egg <- grep("egg", levels(xx$Lifestage), ignore.case = TRUE, value = TRUE)
xx <- within(xx, IsEgg <- Lifestage%in%is.egg)
is.scale <- grep("OS", unique(xx$SLS), ignore.case = TRUE, value = TRUE)
xx <- within(xx, IsScale <- SLS%in%is.scale)
xx <- xx[!is.na(xx$Total), ] # won't total unless

### Define what is dead
xx <- within(xx, dead <- Dead) # for eggs will have some erroneously entered as 0
xx <- within(xx, dead[IsEgg] <- Unhatched[IsEgg]) # overwrites those errors also
xx <- within(xx, Dead[IsScale] <- Dead[IsScale] + Moribund[IsScale])
xx$Row <- seq(nrow(xx))

xxx <- xx %>%
  arrange(Date, SLS, Fruit, Temperature, Duration, Rep, Efpcc) %>%
  select(Date, SLS, Fruit, Temperature, Duration, Rep, Efnom, Efpcc, HC, dead, Total, Row)
w)
idcols <- names(xxx %>%
  select(Date, SLS, Fruit, Temperature, Duration, Rep))
respcols <- names(xxx %>%
  select(dead, Total, Row))

xxx <- within(xxx, Ndx <- paste(Date, SLS,Fruit, Temperature, Duration, Rep, sep = "|"))

### Which are the controls' rows
xx.hc <- xxx[xxx$HC, ] # i.e. handling controls
xx.co2c <- xxx[with(xxx, Efpcc == 0 & !HC),] # i.e. CO2 controls
cont.rows <- rbind(xx.hc, xx.co2c)$Row
treat.rows <- xxx$Row[!xxx$Row %in% cont.rows] # i.e. rows that have treatments applied
co2cIndx <- with(xx.co2c, Ndx)
hcIndx <- with(xx.hc, Ndx)
treatIndx <- unique(xxx$Ndx) # one for every treatment combination
xx.treat <- xxx[xxx$Row %in% treat.rows,]

### Align controls with the corresponding treated data
nocont.df <- NULL
contonly <- NULL
cont.df <- NULL # collect all control data
for(i in treatIndx){
## browser()
hand.i <- xx.hc[xx.hc$Ndx == i,]
co2.i <- xx.co2c[xx.co2c$Ndx == i,]
treat.i <- xx.treat[xx.treat$Ndx == i,]
cont.i <- NULL
if(nrow(treat.i) < 1){
  cat(i, "has no treatment data\n")
  contonly <- c(contonly, i)
}
}

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    } else {
      ## check if any combinations have not controls
      if(nrow(hand.i) == 0){
        hand.i <- treat.i[1, ]
        hand.i[, respcols] <- NA
        hand.i[, c("Efpc", "HC")] <- c(0, 1)
      }
      cont.i <- rbind(hand.i) # get controls back together
      if(nrow(co2.i) == 0){
        co2.i <- treat.i[1, ]
        co2.i[, respcols] <- NA
        co2.i[, c("Efpc", "HC")] <- c(0, 0)
      }
      cont.i <- rbind(cont.i, co2.i) # get controls back together
      cont.i <- within(cont.i, HC <- as.logical(HC)) # coerced to numeric above
      nocont.i <- rbind(hand.i, co2.i, treat.i)
      nocont.i <- within(nocont.i, HC <- as.logical(HC))
      cont.df <- rbind(cont.df, ditch("Ndx", cont.i)) # don't need Ndx
      nocont.df <- rbind(nocont.df, ditch("Ndx", nocont.i)) # don't need Ndx
    }
  }
}

### Get 3 dataframes into one Excel file
contronly.df <- ditch("Ndx", xxx[xxx$Ndx %in% contronly,])
contrmissing.df <- nocont.df[is.na(nocont.df$Row),]

## require("WriteXLS")
## WriteXLS(c("cont.df", "contrmissing.df", "contronly.df"), "LBAMmissing.xls",
##          c("AllControls", "NoControls", "ControlsOnly"))

### Use Duration 3 controls when Duration 2 is without
## put Ndx back in (slightly different one)
contrmissing.df <- within(contrmissing.df,
                          Ndx <- paste(Date, HC, SLS, Fruit, Temperature, Duration, Rep,
sep = "|"))
cont.df <- rbind(cont.df, contronly.df)
cont.df <- within(cont.df,
                  Ndx <- paste(Date, HC, SLS, Fruit, Temperature, Duration, Rep, sep = "|
"))
## browser()
contrmissing.dfb4 <- contrmissing.df
reused <- 0
for(k in seq(nrow(contrmissing.df))){
  missing.k <- contrmissing.df[k, ]
  browser()
  if(is.na(missing.k$Total)){ # otherwise nothing needed
    if(missing.k$Duration == 2){
      Ndx.k <- missing.k$Ndx
      Ndx.kFix <- gsub("\\|2\\|", "|3|", Ndx.k)
      ## if(Ndx.k == "2015-05-29|TRUE|LBAM egg|Kiwifruit|5|2|1")
      mort.dat <- c("dead", "Total")
      ##
      reuse.k <- cont.df[cont.df$Ndx == Ndx.kFix, ]
      if(nrow(reuse.k) > 0){
        contrmissing.df[contrmissing.df$Ndx == Ndx.k, mort.dat] <- reuse.k[, mort.dat]
        reused <- reused + 1
        cat("Reused", Ndx.kFix, "\n")
      }
      ## try(contrmissing.df[contrmissing.df$Ndx == Ndx.k, mort.dat] <-
      ## cont.df[cont.df$Ndx == Ndx.kFix, mort.dat])
    }
  }
}

}

## remove contronly.df Rows from repaired control data (useful ones already copied)
fixed.cont.df <- cont.df[!cont.df$Row %in% contronly.df$Row, ]
cont.notmissing.df <- contrmissing.df[!with(contrmissing.df, is.na(Total)),]
use.df <- ditch("Ndx", rbind(fixed.cont.df, cont.notmissing.df, xx.treat)) %>%
  arrange(Date, SLS, Fruit, Temperature, Duration, Rep, Efnom, Efpc, HC) %>

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%
      filter(!is.na(Total))
use.df <- within(use.df,
      Ndx <- paste(Date, SLS, Fruit, Temperature, Duration, Rep, sep = "|"))
## if(FALSE){

  WriteXLS(c("fixed.cont.df", "contrmissing.dfB4", "contronly.df", "use.df"), out.xls,
    c("AllControls", "NoControls", "ControlsOnly", "ReadyToUse"), BoldHeaderRow =
TRUE,
    FreezeRow = 4, FreezeCol = 3)
## }

## Check if there's any difference between "Controls"
if(FALSE){
  test.control2 <- function(dff){
    dff <- dff[dff$Efpc == 0,] # i.e. controls
    sls <- unique(dff$Ndx)
    cont.out.df <- data.frame(Index = sls)
    cont.out.df <- within(cont.out.df, HC <- CO2 <- Psame <- NA)

    for(sl in sls){
      dfs <- dff[dff$Ndx == sl,]
      cat("\n", sl, ":\n =====\n")
      if(nrow(dfs) == 2){
        spec.glm <- glm(cbind(dead, Total - dead) ~ HC, data = dfs,
          family = binomial)
        hand.mort <- with(dfs[df$HC, ], round(100 * dead/Total))
        CO.mort <- with(dfs[!df$HC, ], round(100 * dead/Total))
        ## browser()
        Psl <- anova(spec.glm, test = "Chi")[2, "Pr(>Chi)"]
        cont.out.df <- within(cont.out.df, Psame[Index == sl] <- Psl)
        cont.out.df <- within(cont.out.df, HC[Index == sl] <- hand.mort)
        cont.out.df <- within(cont.out.df, CO2[Index == sl] <- CO.mort)
      }
    }
    cont.out.df
  }

  aa <- test.control2(use.df)
  browser()
  aa[with(aa, HC > CO2),] # shows about 1/3 have HC > CO2
}
use.df
}

```