

# Appendix0\_10

```
# NDNS analysis, data management -----

# Change the data path accordingly -----
setwd("/home/wangcc-me/Downloads/UKDA-6533-stata11_se/stata11_se/") # on Ubuntu
library(epiDisplay)
library(plyr)
library(tidyverse)

# Read the data into memory -----
library(haven)
data <- read_dta("ndns_rp_yr1-4a_foodleveldietarydata_uk.dta")
data56 <- read_dta("ndns_rp_yr5-6a_foodleveldietarydata.dta")
data78 <- read_dta("ndns_rp_yr7-8a_foodleveldietarydata.dta")
names(data)
names(data56)
names(data78)
names(data)[names(data) == "seriali"] <- "id"
names(data56)[names(data56) == "seriali"] <- "id"
names(data78)[names(data78) == "seriali"] <- "id"

# Extract the data we needed -----
df14d <- data[, c(113, 1, 2, 3, 5, 6, 7, 8, 9, 21, 24, 55, 57, 58,
  59, 60, 61, 62, 63, 64)]
var <- names(df14d)
df56d <- data56 %>% select(var)
df78d <- data78 %>% select(var)
dfs1 <- rbind(df14d, df56d, df78d)
dfs2 <- dfs1[dfs1$Age >= 19, ]
rm(data, data56, data78)
dfs2

# Calculate the time (minute and hour) when they eat -----

dfs2$MealTime_chr <- as.character(dfs2$MealTime)
dfs2$MealTime_hm <- unlist(strsplit(dfs2$MealTime_chr, " "))[c(FALSE,
  TRUE)]
dfs2$MealHourN <- as.numeric(unlist(strsplit(dfs2$MealTime_hm, ":"))[c(TRUE,
  FALSE, FALSE)])
dfs2$MealMinN <- as.numeric(unlist(strsplit(dfs2$MealTime_hm, ":"))[c(FALSE,
  TRUE, FALSE)])
dfs2$MealMinN0 <- (60 * dfs2$MealHourN) + dfs2$MealMinN
dfs3 <- dfs2[order(dfs2$id, dfs2$DayNo, dfs2$MealMinN0), ]
length(unique(dfs3$id)) ## number of participants = 6155

# Create a subset data with only the first observation of each
# participant -----
NDNS <- dfs3[!duplicated(dfs3$id), ]
with(NDNS, tab1(SurveyYear, graph = FALSE, decimal = 2))

# #SurveyYear :
```

	#	Frequency	Percent	Cum. percent
# NDNS Year 1	801	13.01	13.01	
# NDNS Year 2	812	13.19	26.21	
# NDNS Year 3	782	12.71	38.91	
# NDNS Year 4	1055	17.14	56.05	
# NDNS Year 5	625	10.15	66.21	
# NDNS Year 6	663	10.77	76.98	
# NDNS Year 7	703	11.42	88.40	
# NDNS Year 8	714	11.60	100.00	
# Total	6155	100.00	100.00	

```
# create a variable combine id and day No -----
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```
dfs3 <- dfs3 %>%
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```
mutate(id_dy = paste(id, DayNo, sep = "D"))
```

```
# For each subject, the total energy/carbohydrate intake for each eating  
# time can be calculated -----
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```
old <- Sys.time()
```

```
Energy <- ddply(dfs3, .(id_dy, id, SurveyYear, DayNo, Age, Sex,  
                      DiaryDaysCompleted, MealHourN, DayofWeek),  
               summarise,  
                 Tot_Energ = sum(EnergykJ),  
                 Tot_Carb = sum(Carbohydrateg),  
                 Tot_Sugar = sum(Totalsugarsg),  
                 Tot_Starch = sum(Starchg))
```

```
new <- Sys.time() - old
```

```
print(new)
```

```
# Time difference of 3.876385 mins
```

```
rm(df14d, df56d, df78d, dfs2)
```

```
# Calculate the energy from total carbohydrates -----
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```
Energy <- Energy %>%
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```
mutate(KJcarbo = Tot_Carb * 16) %>%
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```
mutate(CarKJpercentage = KJcarbo/Tot_Energ) %>%
```

```
mutate(Carbo = cut(CarKJpercentage, breaks = c(0, 0.26, 0.75, 2),  
              right = FALSE)) %>% mutate(Carbo2 = cut(CarKJpercentage, breaks = c(0,  
              0.26, 2), right = FALSE))
```

```
Energy0 <- Energy[!(Energy$Tot_Energ == 0), ]
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```
# some food consumption does not contain any carbohydrates
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```
Energy0$Carbo <- factor(Energy0$Carbo, labels = c("Low_carb", "Med_carb",  
          "High_carb"))
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Energy0$Carbo2 <- factor(Energy0$Carbo2, labels = c("Low_carb", "Med_or_high_carb"))
```

```
# Generate data sets for each day -----
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```
dta_day1 <- Energy0 %>%
```

```
filter(DayNo == 1) %>%
```

```
select(c("id", "Age",  
          "Sex", "DayofWeek", "MealHourN", "Carbo", "Carbo2")) %>%
```

```
mutate(DayofWeek = factor(DayofWeek,  
              levels = c("Monday", "Tuesday", "Wednesday", "Thursday", "Friday",  
              "Saturday", "Sunday")))
```

```

dta_day2 <- Energy0 %>%
  filter(DayNo == 2) %>%
  select(c("id", "Age",
           "Sex", "DayofWeek", "MealHourN", "Carbo", "Carbo2")) %>%
  mutate(DayofWeek = factor(DayofWeek,
                             levels = c("Monday", "Tuesday", "Wednesday", "Thursday", "Friday",
                                           "Saturday", "Sunday")))

dta_day3 <- Energy0 %>%
  filter(DayNo == 3) %>%
  select(c("id", "Age",
           "Sex", "DayofWeek", "MealHourN", "Carbo", "Carbo2")) %>%
  mutate(DayofWeek = factor(DayofWeek,
                             levels = c("Monday", "Tuesday", "Wednesday", "Thursday", "Friday",
                                           "Saturday", "Sunday")))

dta_day4 <- Energy0 %>%
  filter(DayNo == 4) %>%
  select(c("id", "Age",
           "Sex", "DayofWeek", "MealHourN", "Carbo", "Carbo2")) %>%
  mutate(DayofWeek = factor(DayofWeek,
                             levels = c("Monday", "Tuesday", "Wednesday", "Thursday", "Friday",
                                           "Saturday", "Sunday")))

vecid1 <- unique(dta_day1$id) # n = 6153
vecid2 <- unique(dta_day2$id) # n = 6153
vecid3 <- unique(dta_day3$id) # n = 6151
vecid4 <- unique(dta_day4$id) # n = 6026

Noday1 <- setdiff(vecid, vecid1) # two subjects did not have day 1 data
Noday2 <- setdiff(vecid, vecid2) # two subjects did not have day 2 data
Noday3 <- setdiff(vecid, vecid3) # four subjects did not have day 3 data
Noday4 <- setdiff(vecid, vecid4) # 129 subjects did not have day 4 data

# Transform the data shape from long to wide -----
dta_d1_wide <- dta_day1[, -7] %>%
  spread(key = MealHourN, value = Carbo)
names(dta_d1_wide)[5:28] <- paste(rep("H", 24), 0:23, sep = "")

dta_d2_wide <- dta_day2[, -7] %>%
  spread(key = MealHourN, value = Carbo)
names(dta_d2_wide)[5:28] <- paste(rep("H", 24), 0:23, sep = "")

dta_d3_wide <- dta_day3[, -7] %>%
  spread(key = MealHourN, value = Carbo)
names(dta_d3_wide)[5:28] <- paste(rep("H", 24), 0:23, sep = "")

dta_d4_wide <- dta_day4[, -7] %>%
  spread(key = MealHourN, value = Carbo)
names(dta_d4_wide)[5:28] <- paste(rep("H", 24), 0:23, sep = "")

# recode NA to not eating -----
for (i in 5:ncol(dta_d1_wide))
  if (is.factor(dta_d1_wide[, i])) levels(dta_d1_wide[,

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i]) <- c(levels(dta_d1_wide[, i]), "Not_eating")

dta_d1_wide[is.na(dta_d1_wide)] <- "Not_eating"

for (i in 5:ncol(dta_d2_wide))
  if (is.factor(dta_d2_wide[, i])) levels(dta_d2_wide[,
    i]) <- c(levels(dta_d2_wide[, i]), "Not_eating")

dta_d2_wide[is.na(dta_d2_wide)] <- "Not_eating"

for (i in 5:ncol(dta_d3_wide))
  if (is.factor(dta_d3_wide[, i])) levels(dta_d3_wide[,
    i]) <- c(levels(dta_d3_wide[, i]), "Not_eating")

dta_d3_wide[is.na(dta_d3_wide)] <- "Not_eating"

for (i in 5:ncol(dta_d4_wide))
  if (is.factor(dta_d4_wide[, i])) levels(dta_d4_wide[,
    i]) <- c(levels(dta_d4_wide[, i]), "Not_eating")

dta_d4_wide[is.na(dta_d4_wide)] <- "Not_eating"

```