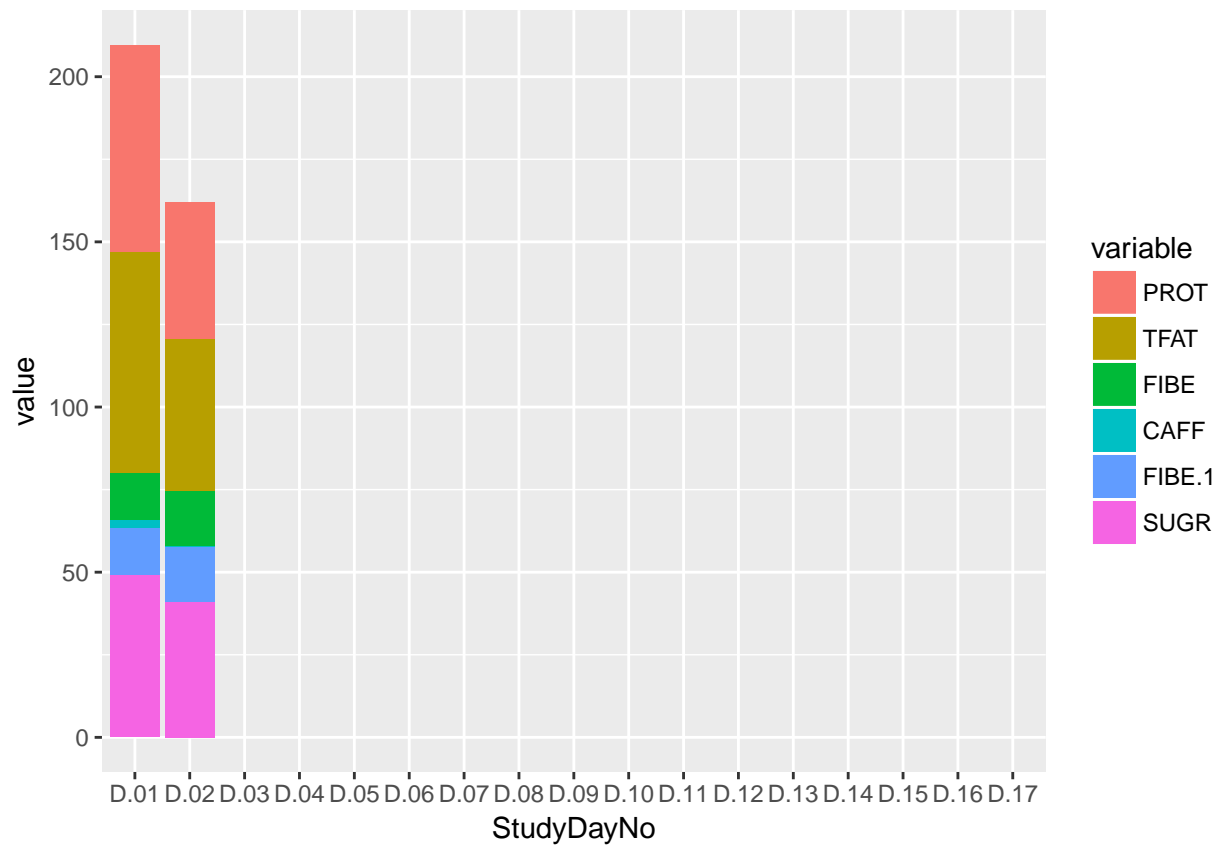


# Subject\_30

Type	Your Average	Total Average
CALORIES	NA	2080.11
PROTEIN	NA	88.57
TOTAL FAT	NA	89.97
CARBS	NA	225.55
FIBER	NA	21.96

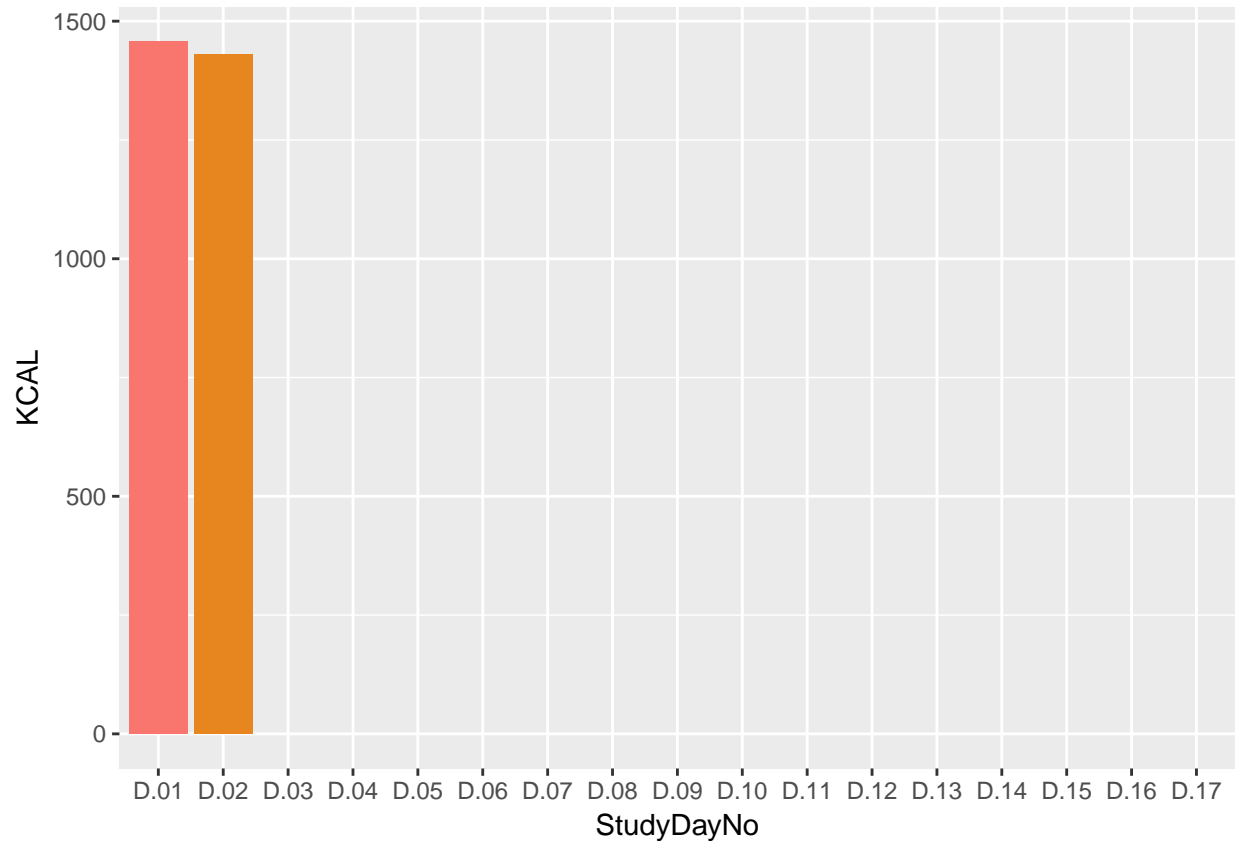
## MicroNutrients

## Warning: Removed 90 rows containing missing values (position\_stack).



## Daily Calorie Intake

## Warning: Removed 15 rows containing missing values (geom\_bar).



## Microbiome Daily Relative Abundance

*# make ggplot bar chart of top 10 most abundant species per day*

```
ggplot(mergedf2, aes(x = StudyDayNo, y = value, fill = rn)) +
  geom_bar(stat = "identity") +
  scale_x_discrete(drop = FALSE) +
  theme_classic() +
  theme(strip.text.y = element_text(angle = 0, size = 8, face = "italic"),
        axis.text.x = element_text(angle = 45, hjust = 1),
        axis.title.x = element_blank(),
        plot.title = element_text(hjust = 0.5),
        strip.background = element_rect(color = "grey")) +
  guides(fill = guide_legend(reverse = TRUE,
                             keywidth = 1,
                             keyheight = 1,
                             ncol = 1)) +
  ylab("Relative Abundance\n") +
  ggtitle("Main species within your gut per day")
```



```

for(i in names(subtaxa)){ dates <- names(subtaxa) #timestamp for each observed sample abund <-
subtaxa[,dates[i]] #abundances for selected timestamps mostabund<- tail(sort(abund),10) #vector of 10
most abundant species (their counts, at least)
}

lst <- list()
for(i in names(subtaxa)){ lst[[i]]<- (subtaxa[,i]) }

```