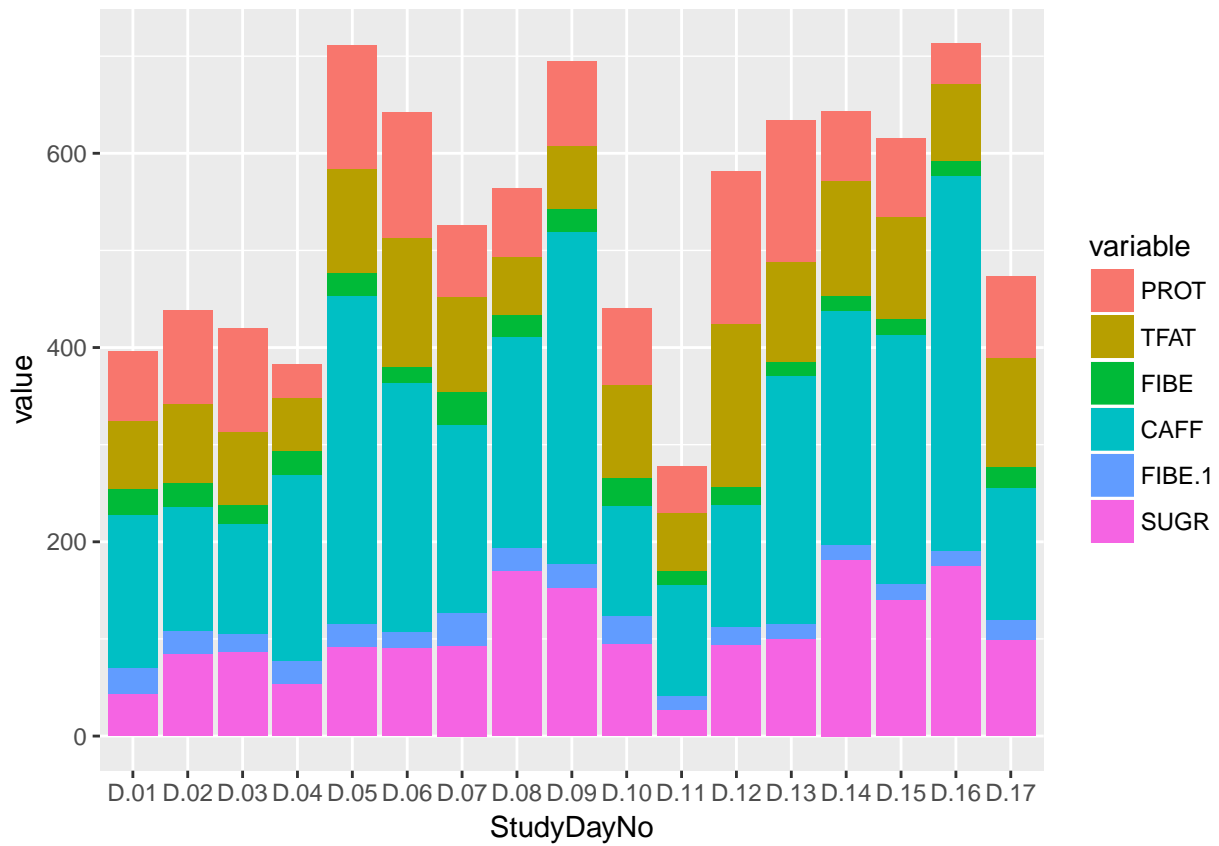


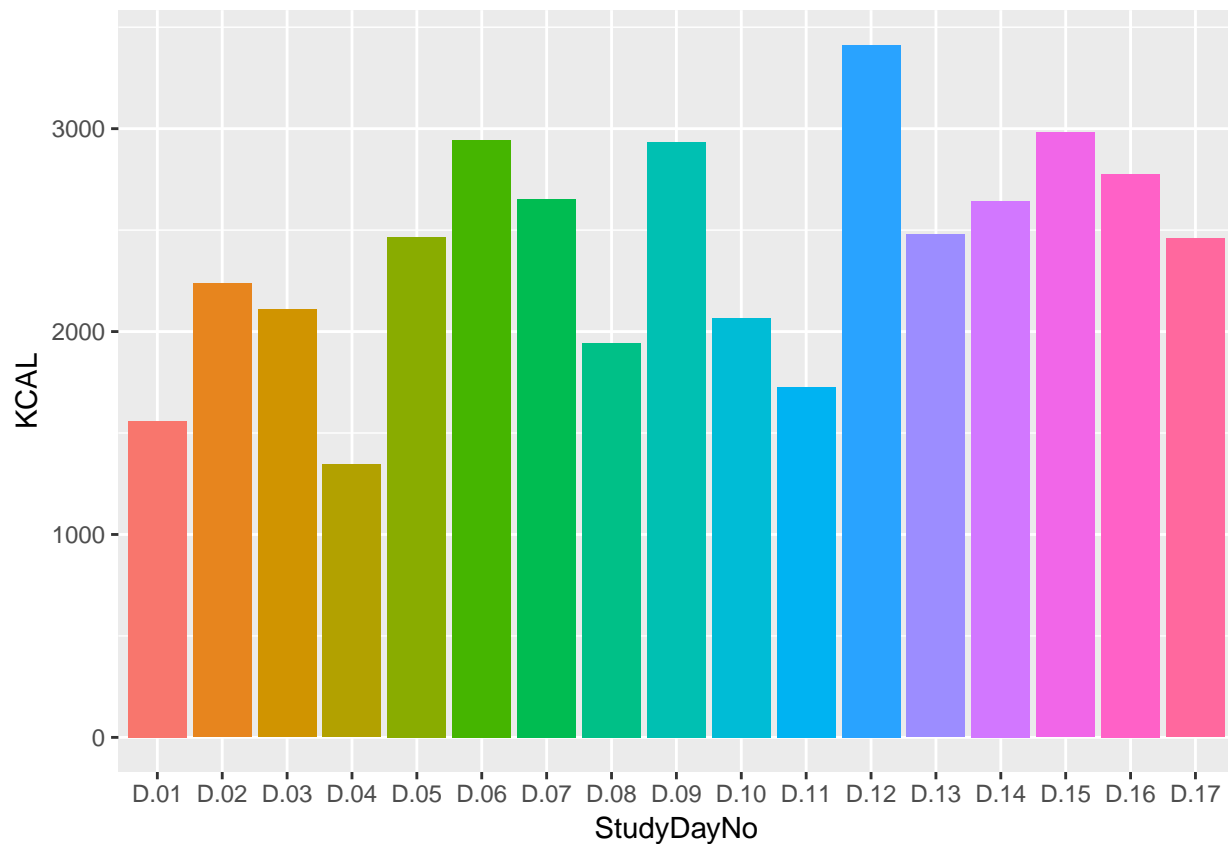
Subject_23

| Type | Your Average | Total Average |
|-----------|--------------|---------------|
| CALORIES | 2396.3 | 2080.11 |
| PROTEIN | 88.52 | 88.57 |
| TOTAL FAT | 93.14 | 89.97 |
| CARBS | 245.34 | 225.55 |
| FIBER | 21.19 | 21.96 |

MicroNutrients



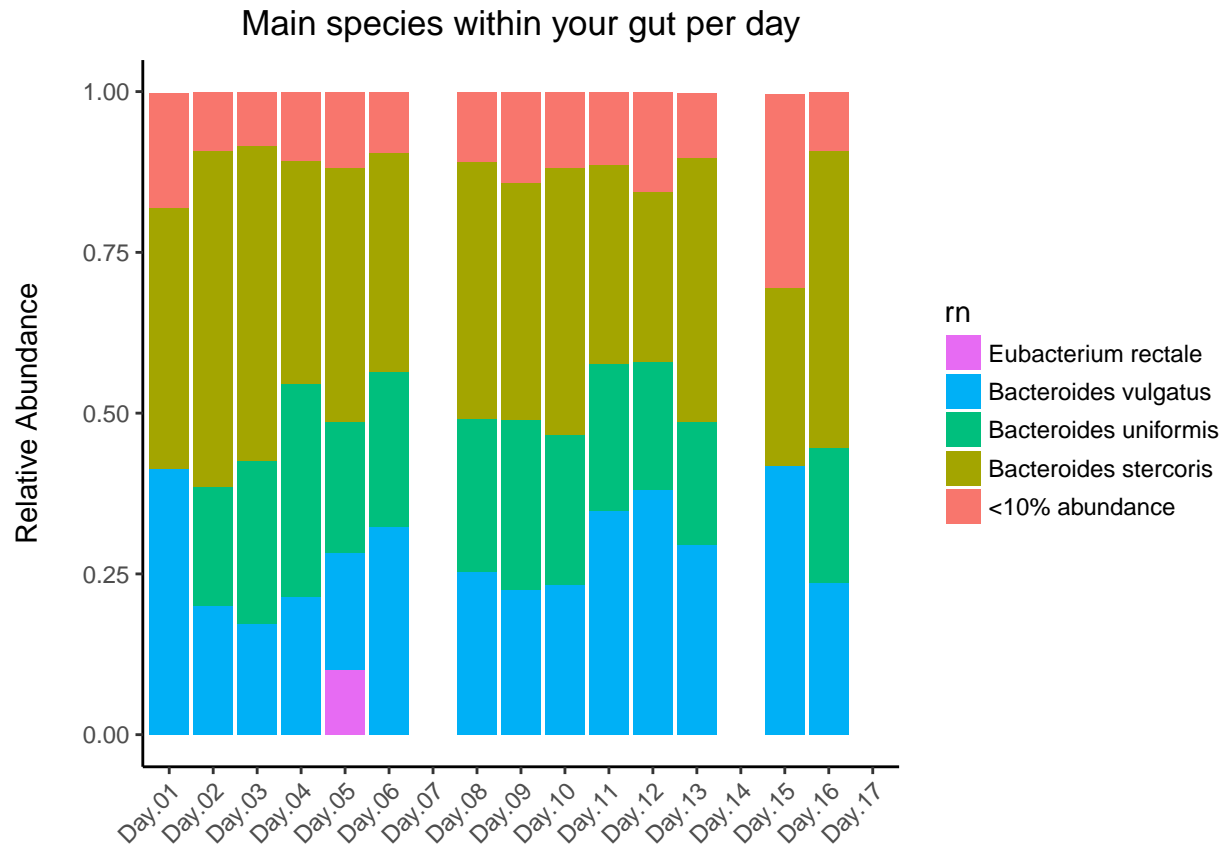
Daily Calorie Intake



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]) }

```