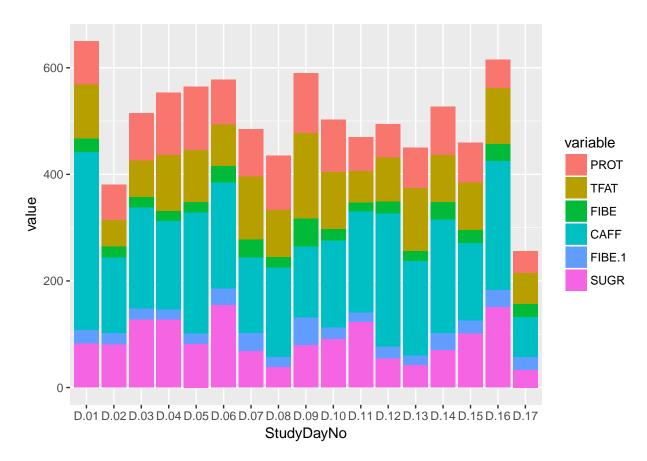
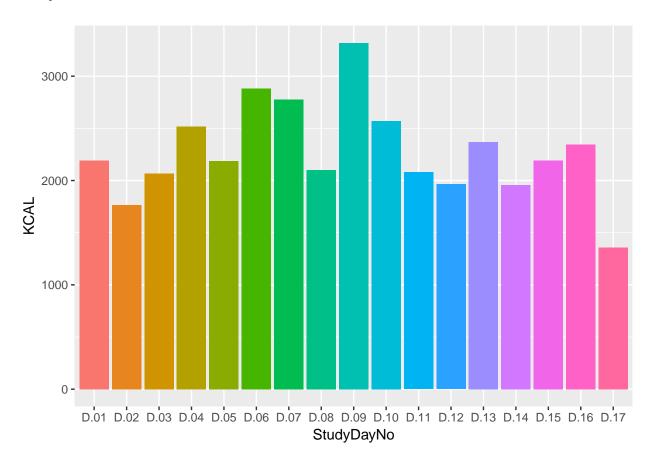
$Subject_41$

| Type | Your Average | Total Average |
|-----------|--------------|---------------|
| CALORIES | 2272.99 | 2080.11 |
| PROTEIN | 83.28 | 88.57 |
| TOTAL FAT | 92.64 | 89.97 |
| CARBS | 262.8 | 225.55 |
| FIBER | 25.41 | 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 most abund<- 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]) } }
```