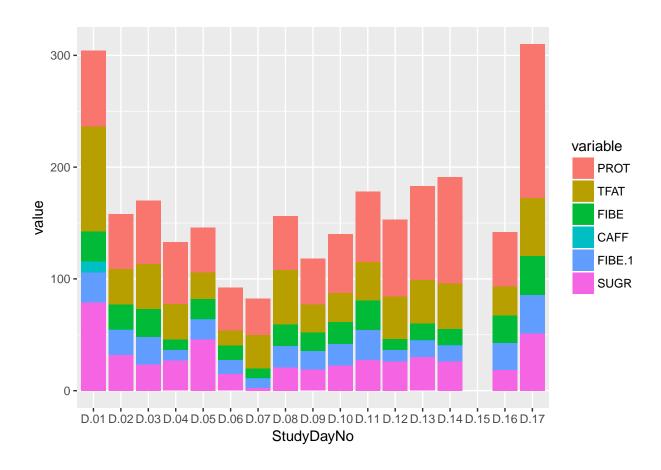
$Subject_28$

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

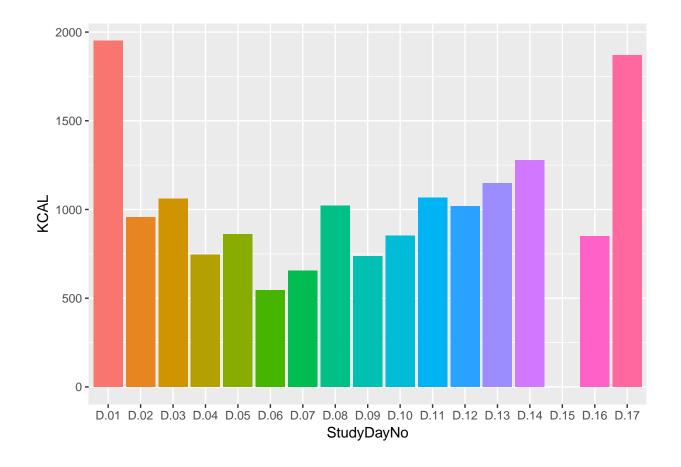
MicroNutrients

Warning: Removed 6 rows containing missing values (position_stack).



Daily Calorie Intake

Warning: Removed 1 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 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]) } }
```