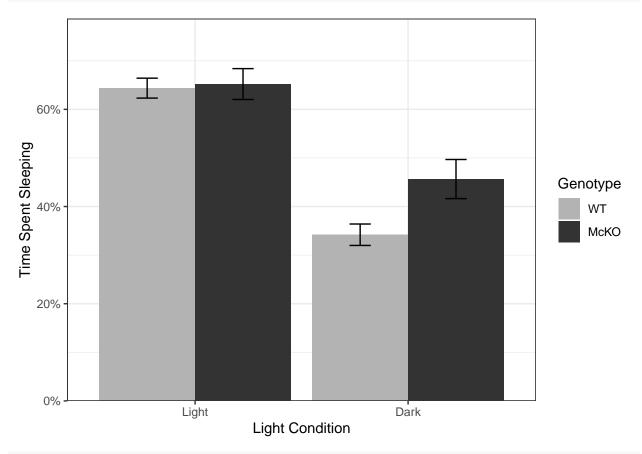
McKO sleep data

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```
library(tidyverse)
## -- Attaching packages ------ tidyverse 1.3.1 --
## v ggplot2 3.3.5
                      v purrr
                                0.3.4
## v tibble 3.1.6
                      v dplyr
                               1.0.7
## v tidyr 1.1.4 v stringr 1.4.0
           2.1.0
## v readr
                      v forcats 0.5.1
## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()
                    masks stats::lag()
library(ggpubr)
#making the dataframe
#from Nik's analysis
Genotype <- c("McKO", "McKO", "WT", "WT")</pre>
condition <- c("Light", "Dark", "Light", "Dark")</pre>
ns \leftarrow c(7, 7, 9, 9)
mean_sleep \leftarrow c(0.6520544, 0.4565667, 0.6436465, 0.3419856)
sds <- c(0.0840014, 0.1064821, 0.0614947, 0.0661068)
ses < c(0.0317495, 0.0402464, 0.0204982, 0.0220356)
sleepdata <- tibble(Genotype, condition, ns, mean_sleep, sds, ses)</pre>
sleepdata <- sleepdata %>%
 mutate(Genotype = fct_relevel(Genotype, c("WT", "McKO")),
        condition = fct_relevel(condition, c("Light", "Dark")))
sleep_plot <- sleepdata %>% ggplot(aes(x=condition, y=mean_sleep, fill=Genotype)) +
 geom_bar(stat="identity", position="dodge") +
 geom_errorbar(stat="identity",
               aes(ymin=mean_sleep-ses, ymax=mean_sleep+ses),
               position=position_dodge(0.9),
               width=0.2) +
 labs(x="Light Condition",
      y="Time Spent Sleeping") +
 scale_y_continuous(labels = scales::percent_format(accuracy = 1),
                    expand = expansion(mult = c(0, 0.15))) +
 scale_fill_manual(values = c("grey70", "grey20")) +
 theme_bw() %+replace%
 theme(plot.background = element_rect(fill = "transparent",colour = NA),
       panel.background = element_rect(fill = "transparent",colour = NA))
```

sleep_plot



ggsave("../figures/sleep_plot.png", plot=sleep_plot, height=3, width=5)