## Plotting Betas

Anton Holm 2020-04-16

```
load("../Rdata_Files/Completed_df.Rda")
load("../Completed_df_2.Rda")
Completed_df_New <- Completed_df %>% full_join(Completed_df_2) %>% mutate(`Random Effects` = replace(`)
## Joining, by = c("simulation", "Limit fraction", "Slope", "Std", "Method", "Intercept", "Beta", "Inte
save(Completed_df_New, file = "../Rdata_Files/Completed_df_New.Rda")
betas <- Completed_df_New %>% filter(Method != "True") %>% mutate(Slope_Perc = ifelse(Slope < 0.01, "1%
tobit_betas_high <- betas %>%
  filter(Method == "Tobit") %>%
  select(-(Method)) %>%
  filter(Std == 1.4) %>%
  filter(`Random Effects` == "High")
museum_betas_high <- betas %>%
  filter(Method == "Museum") %>%
  select(-(Method)) %>%
  filter(Std == 1.4) %>%
  filter(`Random Effects` == "High")
betas_spread_high <- tobit_betas_high %>%
  mutate(Museum_Beta = museum_betas_high$Beta)
Plot_High_Variance <- betas_spread_high %>% ggplot(aes(x=Beta, y=Museum_Beta)) +
  facet_grid(Slope_Perc ~ `Limit fraction`) +
  geom point() +
  geom_hline(data = data.frame(Slope_Perc="1%", y=log(1.01)), mapping=aes(yintercept=y)) +
   geom_vline(data = data.frame(Slope_Perc="1%", x=log(1.01)), mapping=aes(xintercept=x)) +
  geom_hline(data = data.frame(Slope_Perc="5%", y=log(1.05)), mapping=aes(yintercept=y))+
   geom_vline(data = data.frame(Slope_Perc="5%", x=log(1.05)), mapping=aes(xintercept=x)) +
  labs(caption = "Figure 2: Plotting the estimated slopes for the Museum model and Tobit model against
having the variance of both error terms set to high.")
save(Plot_High_Variance, file = "Plot_High_Variance.Rda")
tobit_betas_low <- betas %>%
  filter(Method == "Tobit") %>%
  select(-(Method)) %>%
  filter(Std == 0.05) %>%
```

```
filter(`Random Effects` == "Low")
museum_betas_low <- betas %>%
  filter(Method == "Museum") %>%
  select(-(Method)) %>%
  filter(Std == 0.05) %>%
  filter(`Random Effects` == "Low")
betas_spread_low <- tobit_betas_low %>%
  mutate(Museum_Beta = museum_betas_low$Beta)
Plot_low_Variance <- betas_spread_low %>% ggplot(aes(x=Beta, y=Museum_Beta)) +
  facet_grid(Slope_Perc ~ `Limit fraction`) +
  geom_point() +
   geom_hline(data = data.frame(Slope_Perc="1%", y=log(1.01)), mapping=aes(yintercept=y)) +
   geom_vline(data = data.frame(Slope_Perc="1%", x=log(1.01)), mapping=aes(xintercept=x)) +
  geom_hline(data = data.frame(Slope_Perc="5%", y=log(1.05)), mapping=aes(yintercept=y))+
   geom_vline(data = data.frame(Slope_Perc="5%", x=log(1.05)), mapping=aes(xintercept=x)) +
  labs(caption = "Figure 3: Plotting the estimated slopes for the Museum model and Tobit model against
having the variance of both error terms set to low.")
save(Plot_low_Variance, file = "Plot_low_Variance.Rda")
Plot_low_Variance
```

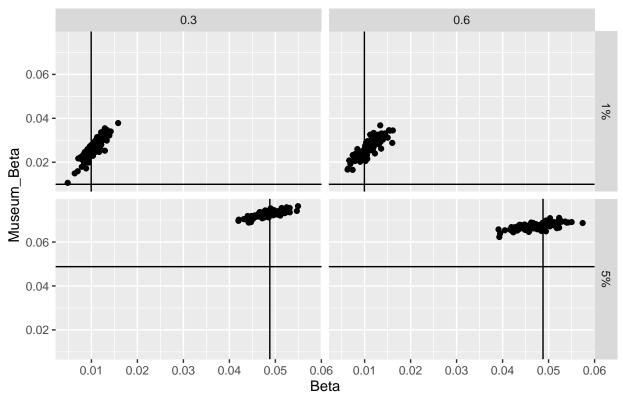


Figure 3: Plotting the estimated slopes for the Museum model and Tobit model against eachother having the variance of both error terms set to low.

Plot\_High\_Variance

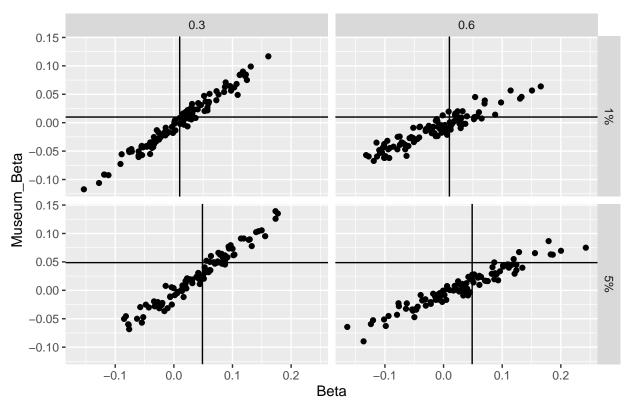


Figure 2: Plotting the estimated slopes for the Museum model and Tobit model against eachother having the variance of both error terms set to high.

```
tobit_betas_individual_high <- betas %>%
  filter(Method == "Tobit") %>%
  select(-(Method)) %>%
  filter(Std == 1.4)
museum_betas_individual_high <- betas %>%
  filter(Method == "Museum") %>%
  select(-(Method)) %>%
  filter(Std == 1.4)
betas_spread_individual_high <- tobit_betas_individual_high %>%
  mutate(Museum_Beta = museum_betas_individual_high$Beta)
Plot_High_Individual_Variance <- betas_spread_individual_high %>% ggplot(aes(x=Beta, y=Museum_Beta, col
  facet_grid(Slope_Perc ~ `Limit fraction`) +
  geom_point(alpha = 0.35) +
   geom_hline(data = data.frame(Slope_Perc="1%", y=log(1.01)), mapping=aes(yintercept=y)) +
   geom_vline(data = data.frame(Slope_Perc="1%", x=log(1.01)), mapping=aes(xintercept=x)) +
  geom_hline(data = data.frame(Slope_Perc="5%", y=log(1.05)), mapping=aes(yintercept=y))+
   geom_vline(data = data.frame(Slope_Perc="5%", x=log(1.05)), mapping=aes(xintercept=x)) +
  labs(caption = "Figure 3: Plotting the estimated slopes for the Museum model
  and Tobit model against eachother having the variance of individual specimens set to high.
  The vertical and horizontal lines correspond to the true value of the slope.")
```

```
save(Plot_High_Individual_Variance, file = "../Rdata_Files/Plot_High_Individual_Variance.Rda")
tobit_betas_individual_low <- betas %>%
  filter(Method == "Tobit") %>%
  select(-(Method)) %>%
  filter(Std == 0.05)
museum_betas_individual_low <- betas %>%
  filter(Method == "Museum") %>%
  select(-(Method)) %>%
  filter(Std == 0.05)
betas_spread_individual_low <- tobit_betas_individual_low %>%
  mutate(Museum_Beta = museum_betas_individual_low$Beta)
Plot_Low_Individual_Variance <- betas_spread_individual_low %>% ggplot(aes(x=Beta, y=Museum_Beta, col =
  facet_grid(Slope_Perc ~ `Limit fraction`) +
  geom_point(alpha = 0.35) +
   geom_hline(data = data.frame(Slope_Perc="1%", y=log(1.01)), mapping=aes(yintercept=y)) +
   geom_vline(data = data.frame(Slope_Perc="1%", x=log(1.01)), mapping=aes(xintercept=x)) +
  geom_hline(data = data.frame(Slope_Perc="5%", y=log(1.05)), mapping=aes(yintercept=y))+
   geom_vline(data = data.frame(Slope_Perc="5%", x=log(1.05)), mapping=aes(xintercept=x)) +
  labs(caption = "Figure 2: Plotting the estimated slopes for the Museum model
  and Tobit model against eachother having the variance of individual specimens set to low.
  The vertical and horizontal lines correspond to the true value of the slope.")
save(Plot_Low_Individual_Variance, file = "../Rdata_Files/Plot_Low_Individual_Variance.Rda")
tobit_betas_individual_medium <- betas %>%
  filter(Method == "Tobit") %>%
  select(-(Method)) %>%
  filter(Std == 0.5)
museum_betas_individual_medium <- betas %>%
  filter(Method == "Museum") %>%
  select(-(Method)) %>%
  filter(Std == 0.5)
betas_spread_individual_medium <- tobit_betas_individual_medium %>%
  mutate(Museum_Beta = museum_betas_individual_medium$Beta)
Plot_Medium_Individual_Variance <- betas_spread_individual_medium %>% ggplot(aes(x=Beta, y=Museum_Beta,
```

```
facet_grid(Slope_Perc ~ `Limit fraction`) +
  geom_point(alpha = 0.35) +
  geom_hline(data = data.frame(Slope_Perc="1%", y=log(1.01)), mapping=aes(yintercept=y)) +
  geom_vline(data = data.frame(Slope_Perc="1%", x=log(1.01)), mapping=aes(xintercept=x)) +
  geom_hline(data = data.frame(Slope_Perc="5%", y=log(1.05)), mapping=aes(yintercept=y))+
  geom_vline(data = data.frame(Slope_Perc="5%", x=log(1.05)), mapping=aes(xintercept=x)) +
  labs(caption = "Figure 3: Plotting the estimated slopes for the Museum model
  and Tobit model against eachother having the variance of individual specimens set to medium.
  The vertical and horizontal lines correspond to the true value of the slope.")

save(Plot_Medium_Individual_Variance, file = "../Rdata_Files/Plot_Medium_Individual_Variance.Rda")
```