

Propensity Score Diagnostics

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2021-09-01 (updated: 2021-10-21)

Checking balance

Love plots (Standardized Mean Difference)

ECDF plots

Standardized Mean Difference (SMD)

$$d = \frac{\bar{x}_{treatment} - \bar{x}_{control}}{\sqrt{\frac{s_{treatment}^2 + s_{control}^2}{2}}}$$

SMD in R

```
library(smd)
library(tidyverse)

df %>%
  # w is optional
  summarise(smd = smd(confounder_1, exposure, w = wts)$estimate)
```

SMD in R

2 Calculate standardized mean differences

```
smds <- df %>%  
  summarise(  
    across(  
      c(confounder_1, confounder_2, ...),  
      list(  
        unweighted = ~smd(.x, exposure)$estimate,  
        weighted = ~smd(.x, exposure, wts)$estimate  
      )  
    )  
  )
```

SMD in R

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SMD in R

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      )  
    )  
  )
```

SMD in R

3 Pivot the SMDs into tidy format

```
plot_df <- smds %>%  
  pivot_longer(  
    everything(),  
    values_to = "SMD",  
    names_to = c("variable", "Method"),  
    names_sep = "_"  
  )
```


SMD in R

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    names_to = c("variable", "Method"),  
    names_sep = "_"  
  )
```

SMD in R

4 Plot them! (in a Love plot!)

```
ggplot(  
  data = plot_df,  
  aes(x = abs(SMD), y = variable, group = Method, color = Method)  
) +  
  geom_line(orientation = "y") +  
  geom_point() +  
  geom_vline(xintercept = 0.1, color = "black", size = 0.1)
```

SMD in R

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SMD in R

4 Plot them! (in a Love plot!)

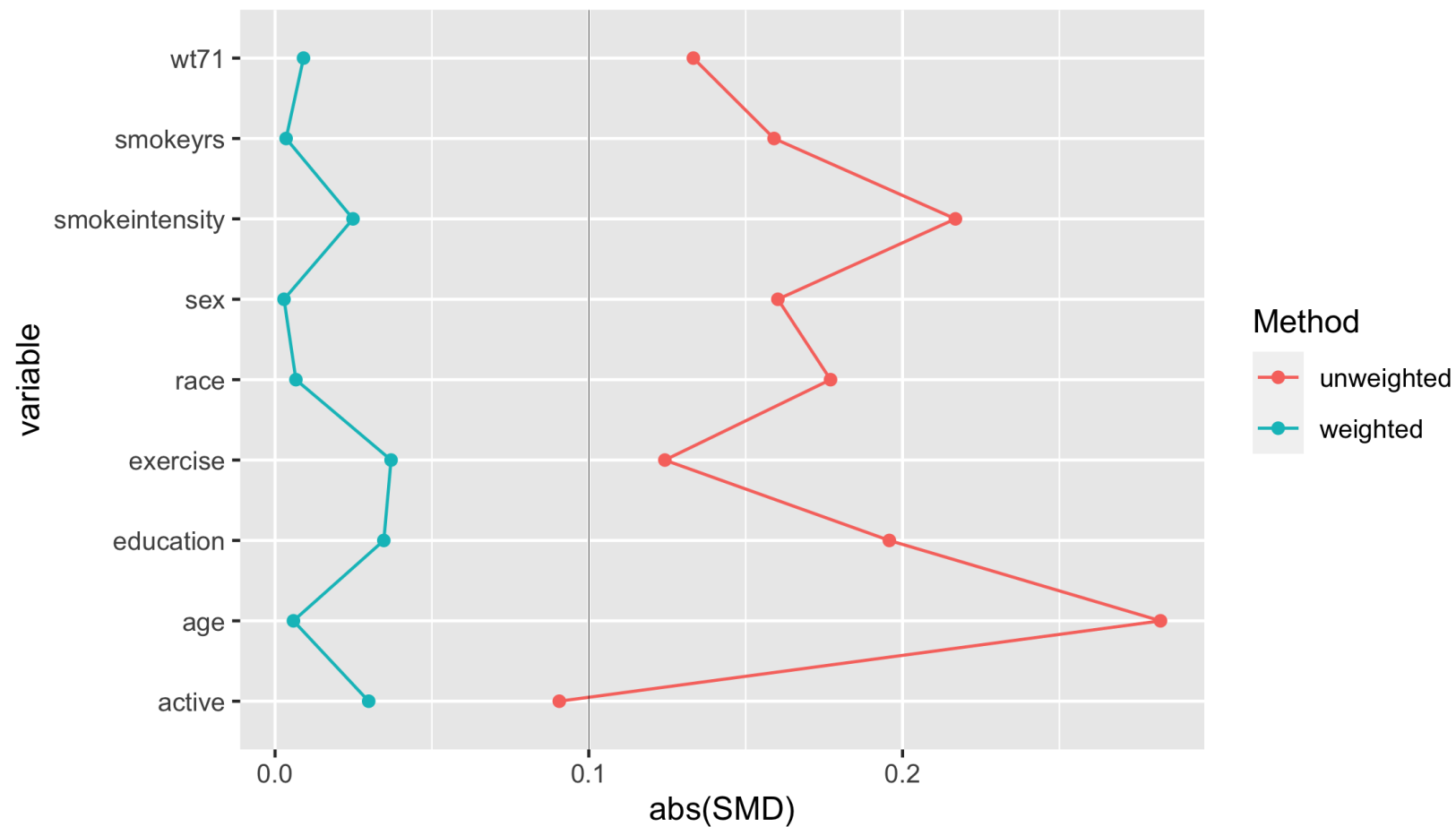
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SMD in R

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```

Love plot



Your turn 1

- 1 **Create a Love Plot for the propensity score weighting you created in the previous exercise**

10:00

ECDF

For continuous variables, it can be helpful to look at the **whole** distribution pre and post-weighting rather than a single summary measure

Unweighted ECDF

```
ggplot(df, aes(x = wt71, group = qsmk, color = factor(qsmk))) +  
  stat_ecdf() +  
  scale_color_manual("Quit smoking", values = c("#5154B8", "#5DB854"),  
                    labels = c("Yes", "No")) +  
  xlab("Weight in Kg in 1971") +  
  ylab("Proportion <= x")
```

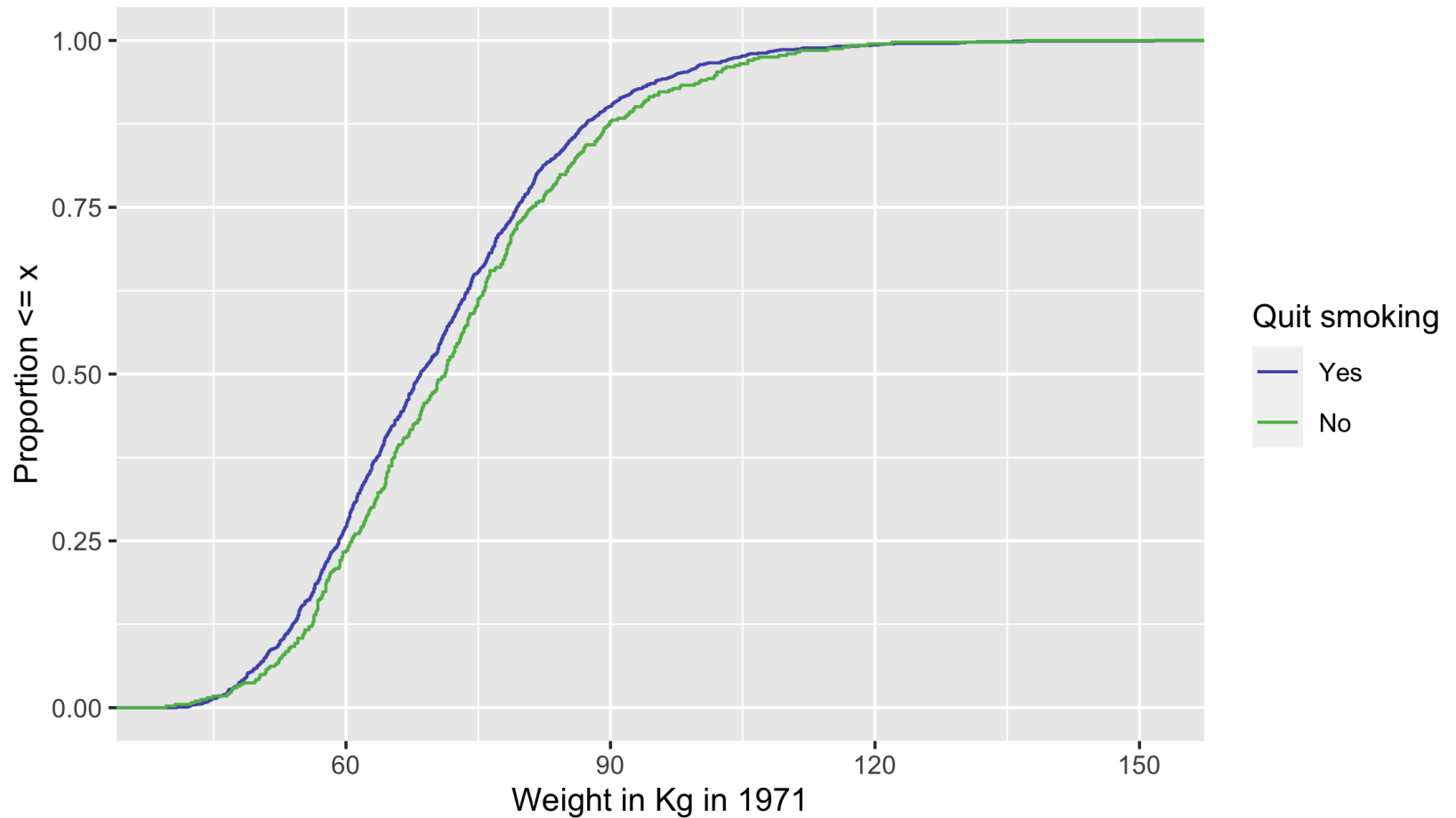
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Unweighted ECDF



Weighted ECDF

```
ecdf_1 <- df %>%  
  filter(qsmk == 1) %>%  
  arrange(wt71) %>%  
  mutate(cum_pct = cumsum(w_ate) / sum(w_ate))  
  
ecdf_0 <- df %>%  
  filter(qsmk == 0) %>%  
  arrange(wt71) %>%  
  mutate(cum_pct = cumsum(w_ate) / sum(w_ate))  
  
ggplot(ecdf_1, aes(x = wt71, y = cum_pct)) +  
  geom_line( color = "#5DB854") +  
  geom_line(data = ecdf_0, aes(x = wt71, y = cum_pct), color = "#515151") +  
  xlab("Weight in Kg in 1971") +  
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Weighted ECDF

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Weighted ECDF

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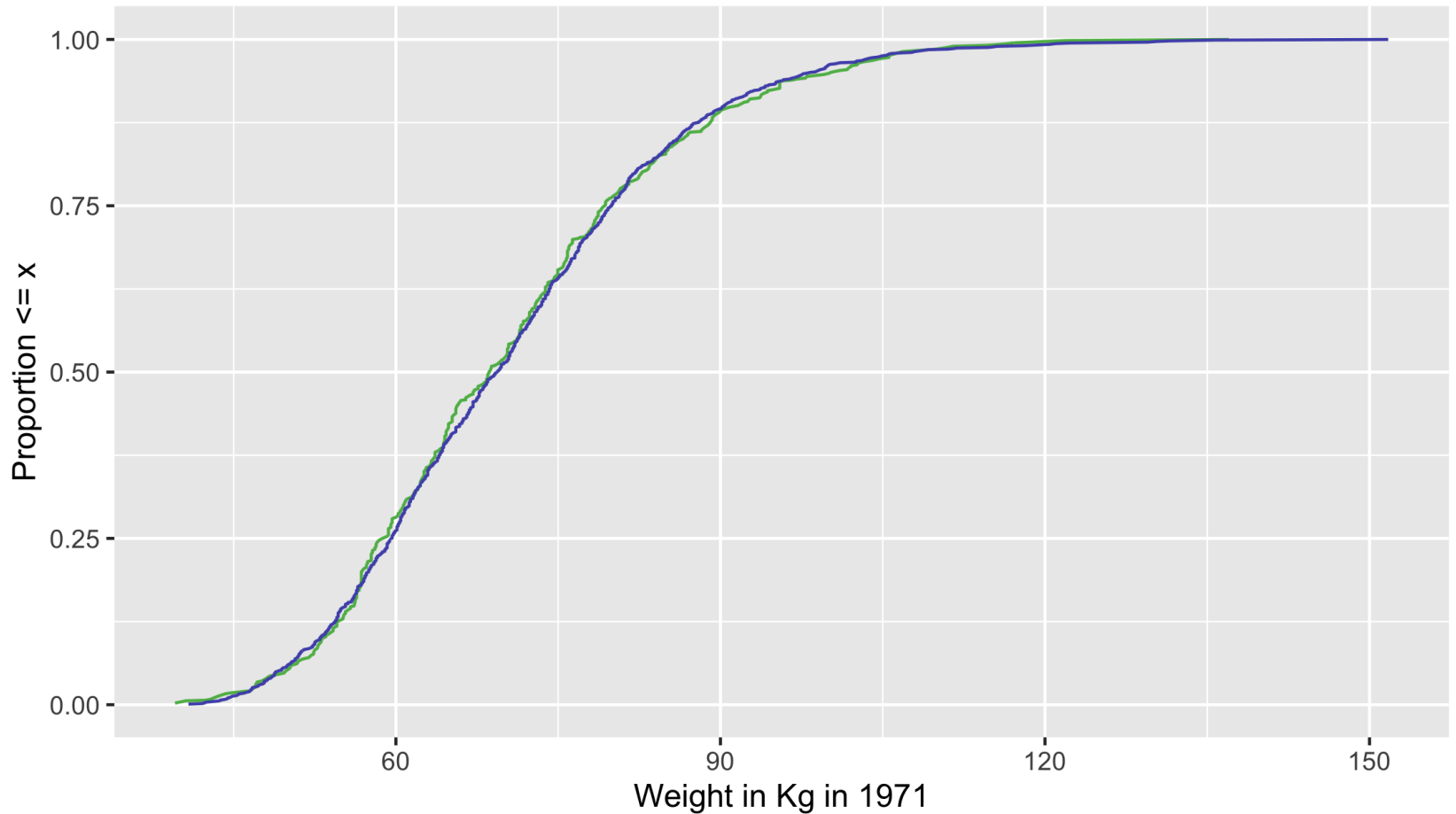
Weighted ECDF

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Weighted ECDF



Your turn 2

- 1 Create an unweighted ECDF examining the smokeyrs confounder for those that quit smoking and those that did not**
- 2 Create a weighted ECDF examining the smokeyrs confounder**

10:00

Bonus! Weighted Tables in R

Weighted Tables in R

1 Create a "design object" to incorporate the weights

```
library(survey)

svy_des <- svydesign(
  ids = ~ 1,
  data = df,
  weights = ~ wts,
  strata = ~ x
)
```

Weighted Tables in R

2 Pass to `gtsummary::tbl_svysummary()`

```
library(gtsummary)
tbl_svysummary(svy_des, by = x) %>%
  add_difference(everything() ~ "smd")
# modify_column_hide(ci) to hide CI column
```

Characteristic	0, N = 1,565 ¹	1, N = 1,561 ¹	Difference ²
WEIGHT IN KILOGRAMS IN 1971	69 (60, 80)	69 (59, 79)	0.01
0: WHITE 1: BLACK OR OTHER IN 1971			0.01
0	1,359 (87%)	1,352 (87%)	
1	206 (13%)	209 (13%)	
AGE IN 1971	43 (33, 52)	43 (33, 53)	-0.01
0: MALE 1: FEMALE			0.00
0	764 (49%)	764 (49%)	
1	802 (51%)	797 (51%)	
NUMBER OF CIGARETTES SMOKED PER DAY IN 1971	20 (10, 25)	20 (10, 30)	0.02
YEARS OF SMOKING	24 (15, 33)	24 (14, 33)	0.00
IN RECREATION, HOW MUCH EXERCISE? IN 1971, 0:much exercise,1:moderate exercise,2:little or no exercise			0.04
0	302 (19%)	294 (19%)	
1	665 (42%)	691 (44%)	
2	599 (38%)	576 (37%)	
IN YOUR USUAL DAY, HOW ACTIVE ARE YOU? IN 1971, 0:very active, 1:moderately active, 2:inactive			0.03
0	700 (45%)	684 (44%)	
1	718 (46%)	738 (47%)	
2	147 (9.4%)	138 (8.9%)	

¹Median (IQR); n (%)

²Standardized Mean Difference