CL Immunization 06

Group K

2025-09-19

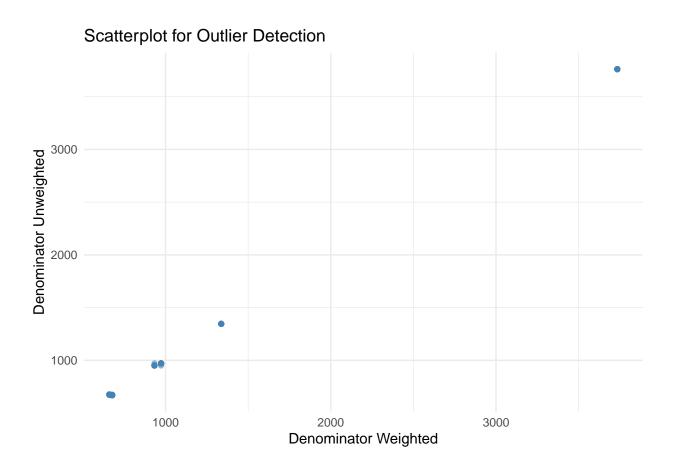
Display Dataset content

```
## # A tibble: 5 x 29
            DataId Indicator Value Precision DHS_CountryCode CountryName SurveyYear
##
     ISO3
##
     <chr> <chr> <chr>
                             <chr> <chr>
                                              <chr>>
                                                               <chr>>
                                                                           <chr>>
## 1 #coun~ #meta~ #indicat~ #ind~ #indicat~ <NA>
                                                               #country+n~ #date+year
## 2 ZAF
            330965 BCG vacc~ 96.8 1
                                              ZA
                                                               South Afri~ 1998
## 3 ZAF
            139796 BCG vacc~ 94.9 1
                                              ZA
                                                               South Afri~ 1998
                                              ZA
## 4 ZAF
            330966 DPT 1 va~ 93.3
                                                               South Afri~ 1998
                                              ZA
                                                              South Afri~ 1998
## 5 ZAF
            139797 DPT 1 va~ 93.1 1
## # i 21 more variables: SurveyId <chr>, IndicatorId <chr>, IndicatorOrder <dbl>,
## #
       IndicatorType <chr>, CharacteristicId <dbl>, CharacteristicOrder <dbl>,
       CharacteristicCategory <chr>, CharacteristicLabel <chr>,
       ByVariableId <chr>, ByVariableLabel <chr>, IsTotal <dbl>,
## #
       IsPreferred <dbl>, SDRID <chr>, RegionId <lgl>, SurveyYearLabel <dbl>,
## #
## #
       SurveyType <chr>, DenominatorWeighted <dbl>, DenominatorUnweighted <dbl>,
       CILow < lgl>, CIHigh < lgl>, LevelRank < lgl>
#Remove the first row(meta data)
imm_df <- imm_df[-1, ]</pre>
```

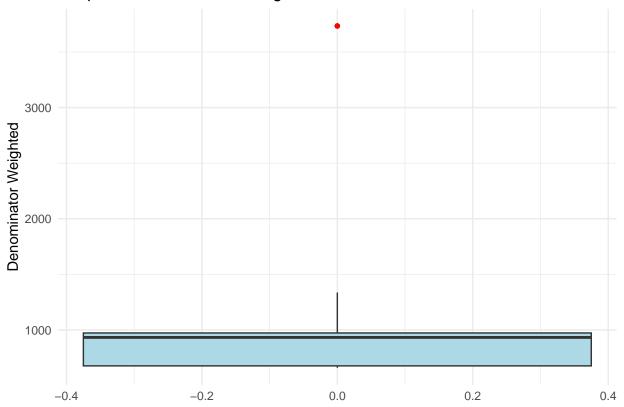
Convert Data Types

#check for unique values

#Drop the countries only one unquie value: reason, there is no useful information - county is also always za #Assumed pattern, the missing values can be filled with the previous non missing value in the opposite attribute







Outlier Handling

```
# Calculate IQR boundaries
Q1_w <- quantile(imm_df$DenominatorWeighted, 0.25, na.rm = TRUE)
Q3_w <- quantile(imm_df$DenominatorWeighted, 0.75, na.rm = TRUE)
IQR_w <- Q3_w - Q1_w
lower_w <- Q1_w - 1.5 * IQR_w
upper_w <- Q3_w + 1.5 * IQR_w
Q1 uw <- quantile(imm df$DenominatorUnweighted, 0.25, na.rm = TRUE)
Q3_uw <- quantile(imm_df$DenominatorUnweighted, 0.75, na.rm = TRUE)
IQR_uw \leftarrow Q3_uw - Q1_uw
lower_uw <- Q1_uw - 1.5 * IQR_uw</pre>
upper_uw <- Q3_uw + 1.5 * IQR_uw
# Cap values to the IQR limits
imm_df <- imm_df %>%
 mutate(
    DenominatorWeighted = pmin(pmax(DenominatorWeighted, lower_w), upper_w),
    DenominatorUnweighted = pmin(pmax(DenominatorUnweighted, lower_uw), upper_uw)
  )
```