

Learning *R*

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Missing values

1. Missing Values

Missing Values - NA's, NaN, Infs

Missing values (NA) can occur

- Data lacking values (are these MCAR, MAR, or IM)?
- Illegal computations ("Carl"/3) or 10/0
- Over/Under flow (e.g. $\exp(1000)$)

Default action of *R* is to propagate missing values, i.e. $3 + \text{NA}$ is also NA.

You can change this action in several ways (the `na.actions`).

R Missing Values

Look at *cereal* dataframe and the *weight* variable.

Some cereals have missing values.

Compare the following results.

Read *help(is.na)* and *help(na.omit)*

```
1 mean(cereal$weight)
2 mean(cereal$weight, na.rm=TRUE)
3 mean(na.omit(cereal$weight))
4
5 length(cereal$weight) # includes missing values
6 length(na.omit(cereal$weight))
7 is.na(cereal$weight)
8 sum( is.na(cereal$weight)) # count num missing
9
10 dim(cereal)
11 dim(na.omit(cereal)) # drop row with missing data
12
13 complete.cases(cereal)
14 dim(cereal[complete.cases(cereal),])
```

Look at cereal dataframe and the *weight* variable.
Compute the % weight of fat, i.e. grams of fat/serving size.
Compute the protein:fat ratio.

```
1 # Using NA in operations leads to NA's
2 cereal$prop.fat <- cereal$fat /cereal$weight
3 cereal$prop.fat
4 is.na(cereal$protein.fat) # Inf is a value
5
6
7 # NA is different from Inf;  protein/fat ratio
8 cereal$protein.fat <- cereal$protein / cereal$fat
9 cereal$protein.fat
10 iis.na(cereal$protein.fat) # Inf is a value
11 is.infinite(cereal$protein.fat)
```

Missing values typically don't show up on plots!

```
1 ggplot(data=cereal, aes(x=weight, y=calories))+  
2   ggtitle("calories vs. Weight per serving")+  
3   xlab("Weight per serving")+ylab("calories")+  
4   geom_jitter()+  
5   geom_smooth(method="lm", se=FALSE)
```

Warning messages:

```
1: Removed 2 rows containing missing values (stat_smooth).  
2: Removed 2 rows containing missing values (geom_point).
```

Look at cereal dataframe and the weight variable.
Regress calories against serving size.

```
1 fit.cal.serving <- lm(calories ~ weight,  
2                       data=cereal)  
3 summary(fit.cal.serving) # note missingness
```

Residual standard error: 12.89 on 73 degrees of freedom
(2 observations deleted due to missingness)

Look at cereal dataframe and the weight variable.
Regress calories against serving size.

```
1 fitted(fit.cal.serving) # only length 75 despite having ind
2 length(fitted(fit.cal.serving))
```

So the following fails:

```
1 cereal$fitted <- fitted(fit.cal.serving)
```

```
Error in '$<-.data.frame'('*tmp*', "fitted", value = c(100.6
  replacement has 75 rows, data has 77
```

Look at cereal dataframe and the weight variable.
Regress calories against serving size.

It is possible to have `lm()` deal *nice*ly with NA, but this approach is not implemented consistently in R.

```
1 fit.cal.serving2 <- lm(calories ~ weight,  
2                       na.action=na.exclude,  
3                       data=cereal)  
4 summary(fit.cal.serving2)  
5 fitted(fit.cal.serving2) # now padded with NA  
6 length(fitted(fit.cal.serving2))  
7 cereal$fitted <- fitted(fit.cal.serving2)
```

Look at cereal dataframe and the weight variable.
Regress calories against serving size.

I prefer to use the predict with new data that propagates missing values properly.

Compare:

```
1 predict(fit.cal.serving)
2 predict(fit.cal.serving, newdata=cereal)
3
4 predict(fit.cal.serving2)
```

- ALWAYS check for NA's in data and ask if MCAR, MAR, or IM

- Sometime special codes are used, e.g. -1 means missing

```
1 mydf [ is.na(mydf$var), ] # list rows where var is missing
2 mydf [ mydf == -1 ] <- NA # common code
```

- NA and Inf's propagate through computation but R is NOT consistent.
 - `mean(var)` is NA if var contains any missing
 - `mean(var, na.rm=TRUE)` drops the NAs
 - `lm(Y ~ X, data=blah)` drops missing rows automatically
- Different results from methods with `na.action=na.exclude` vs. `na.action = na.omit`.