

Let us create a df with some missing cells:

```
col1<-seq(11,21); col2<-seq(23,30); col3<-seq(43,50)
length(col1)<-10; length(col2)<-10; length(col3)<-9
df <-cbind(col1,col2,col3)
```

And inspect for missings:

```
sum(is.na(df))
```

This correctly identifies the number of missing values as 3. We can either remove the na cells to get a complete df by the following:

```
df1<- na.omit(df)
```

Or replace them by column mean using the zoo package:

```
na.aggregate(df)
```

or, to drop columns with all missing values:

```
df <- df[,colSums(is.na(df))<nrow(df)]
```