df-data-case_match

Data

```
library(tidyverse)
df <- data.frame(
  Disease = c(0,0,1,1,0),
  bioMarker = c(31.9,32.0,39.0,32.3,41.0)
)
df
```

```
  Disease bioMarker
1       0      31.9
2       0      32.0
3       1      39.0
4       1      32.3
5       0      41.0
```

Use `case_match` to recode

- replacing `recode function on a vector (not dataframe)`

```
library(tidyverse)
df$Disease <- case_match(
              df$Disease,
              0 ~ "noCancer",
              1 ~ "Cancer"
         )
df
```

```
   Disease bioMarker
1 noCancer      31.9
2 noCancer      32.0
3   Cancer      39.0
4   Cancer      32.3
5 noCancer      41.0
```