

# 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