

```
> data <- read.table("test.txt", header = TRUE)
> data
```

	localisation	tumorsize	progress
XX348	proximal	6.3	FALSE
XX234	distal	8.0	TRUE
XX987	proximal	10.0	FALSE

```
> mode(data)
[1] "list"
> class(data)
[1] "data.frame"
> █
```