```
"patient_id", "age", "gender", "diagnosis", "bmi", "smoker"
"Poo1", 34, "Male", "Cancer", 22.5, "Yes"
"Poo2", 28, "Female", "Normal", 20.5, "No"
"Poo3", 45, "Female", "Cancer", 26.7, "Yes"
"Poo4", 39, "Male", "Normal ", 23.8, "No"
"Poo5", 50, "Female", "Cancer", 27.1, "Yes"
"Poo6", 30, "Male", "Normal", 21.9, "No"
"Poo7", 41, "Female", "Cancer", 25.4, "Yes"
"Poo8", 36, "Female", "Normal", 24.2, "No"
"Poo9", 55, "Male", "Cancer", 28.6, "Yes"
"Po10", 29, "Female", "Normal", 19.8, "No"
"Po11", 48, "Male", "Normal", 24.7, "No"
"Po12", 33, "Female", "Cancer", 22.1, "Yes"
"Po13", 42, "Female", "Cancer", 27.9, "Yes"
"Po14", 31, "Male", "Normal", 23.5, "No"
"Po15", 38, "Female", "Cancer", 26.1, "Yes"
"Po16", 53, "Male", "Cancer", 28, "Yes"
"Po17", 26, "Female", "Normal", 21, "No"
"Po18", 46, "Male", "Cancer", 25.6, "Yes"

# convert height into Factor
data$height_fac <-as.character.factor(data$height)
"Po19", 40, "Female", "Normal", 22.9, "No"
"Po20", 35, "Male", "Cancer", 24.3, "Yes"</pre>
```