demo table\_subtype\_by\_demographics()

library(fxtas)

patient\_data = sim\_data |>  
 dplyr::filter(.data$Category == "Patient")  
table = sim\_subtype\_and\_stage\_table  
tab1 <- table\_subtype\_by\_demographics(  
 patient\_data,   
 table,  
 demographic\_vars = "Sex",  
 footnotes\_as\_letters = FALSE) |>   
 gtsummary::as\_flex\_table(ref\_symbols = letters)  
tab1

|  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
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| Table 1: test   | **Characteristic** | **Overall**  N = 18a | **Subtype 1**  N = 1a | **Subtype 3**  N = 17a | **p-value** | | --- | --- | --- | --- | --- | | **Sex, n (%)** |  |  |  | 1.000b | | Female | 9 (50%) | 1 (100%) | 8 (47%) |  | | Male | 9 (50%) | 0 (0%) | 9 (53%) |  | | an (column %) | | | | | | bGroup comparison was done by Fisher's exact test | | | | | |