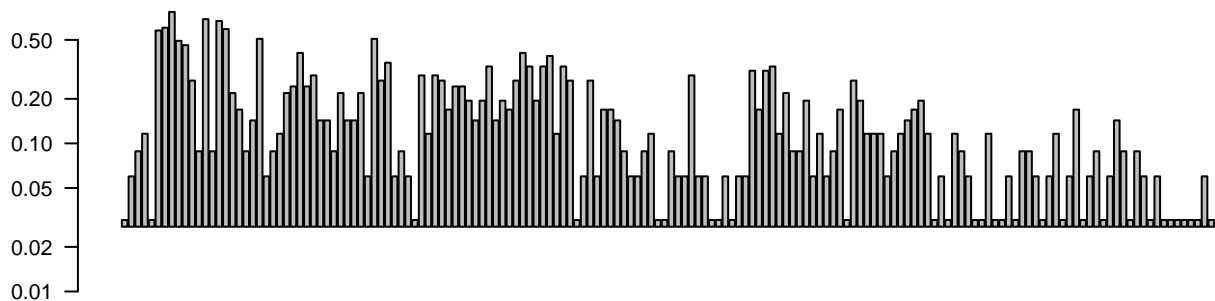


n. mutated genes per sample

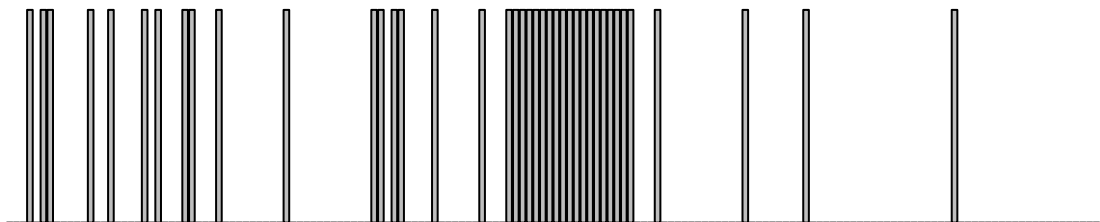


p(one mutation in this pathway)



Expected number of samples with mutation in this pathway = 25

Samples with mutations in this pathway = 40



samples