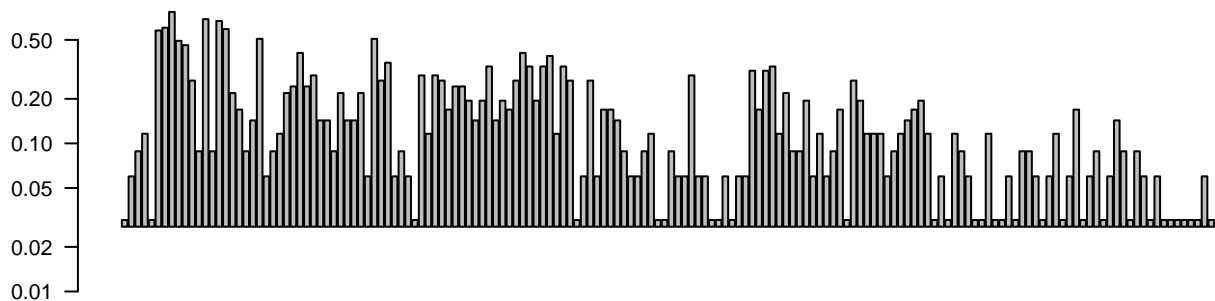


**n. mutated genes per sample**

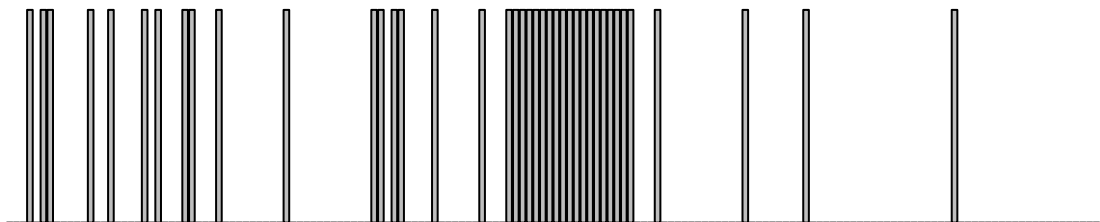


**p(one mutation in this pathway)**



Expected number of samples with mutation in this pathway = 25

**Samples with mutations in this pathway = 40**



samples