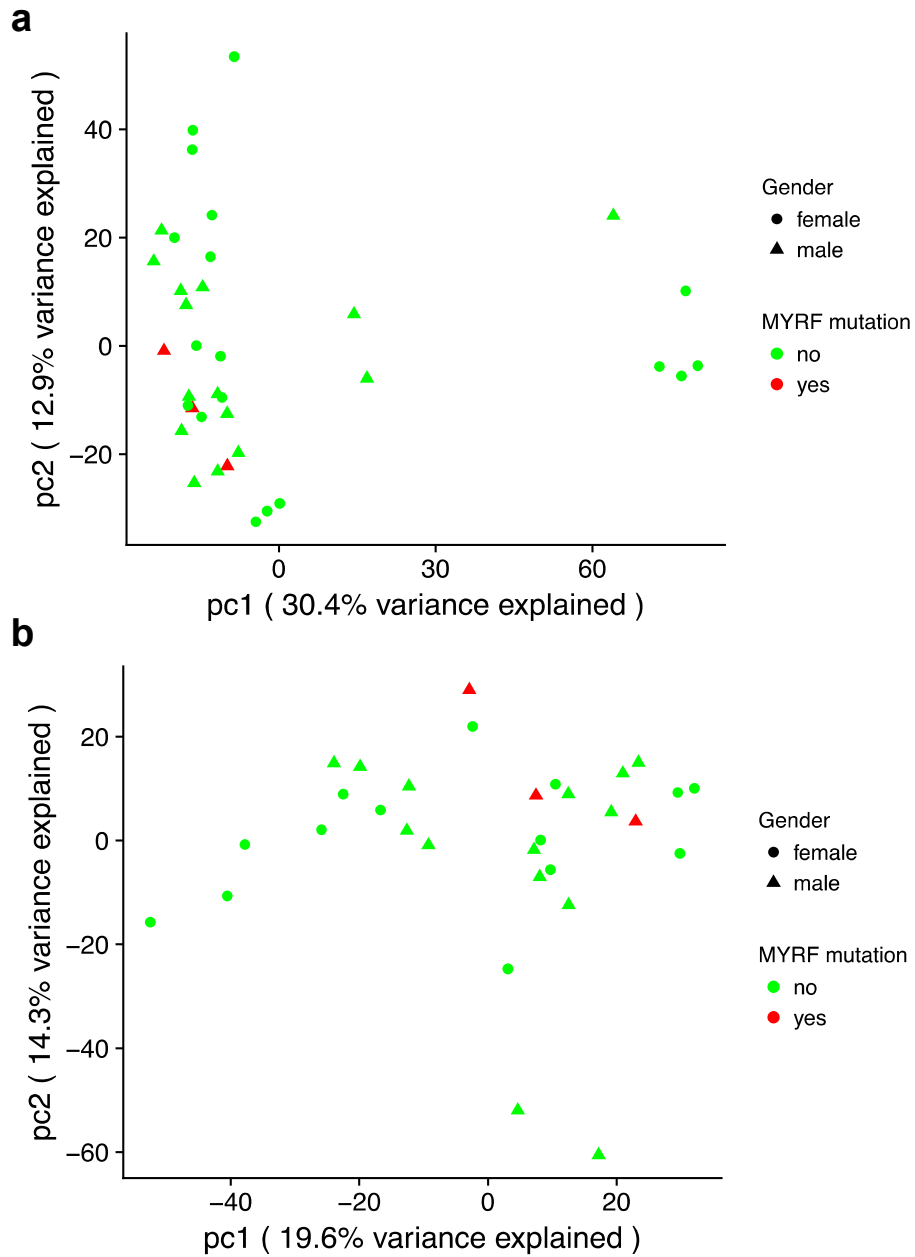


S5 Fig. Principle component (PC) analysis of RNA-seq samples.



(a) The expression profile of five samples were separated from the other samples on the first two PC axis.

(b) After removing outliers, no obvious outlier can be found, and all remaining 31 samples were used in downstream analysis. Three samples from patients with *MYRF* damaging variants are indicated.