

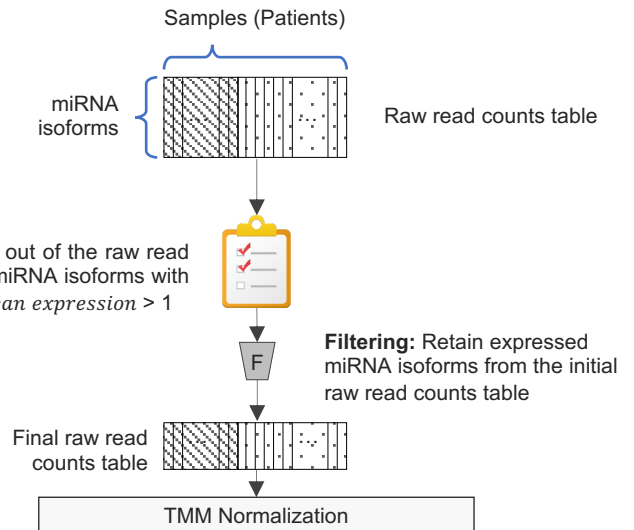


## Input preprocessing

For each race group/cancer tissue (e.g., Black or African American/Solid Primary Tumor)

-  Patients with event (i.e., death or relapse)
-  Patients without event (i.e., alive or no relapse)

Calculate the RPM out of the raw read counts and retain miRNA isoforms with *RPM geometric mean expression* > 1



## Risk Score-based Prognostic Signature

