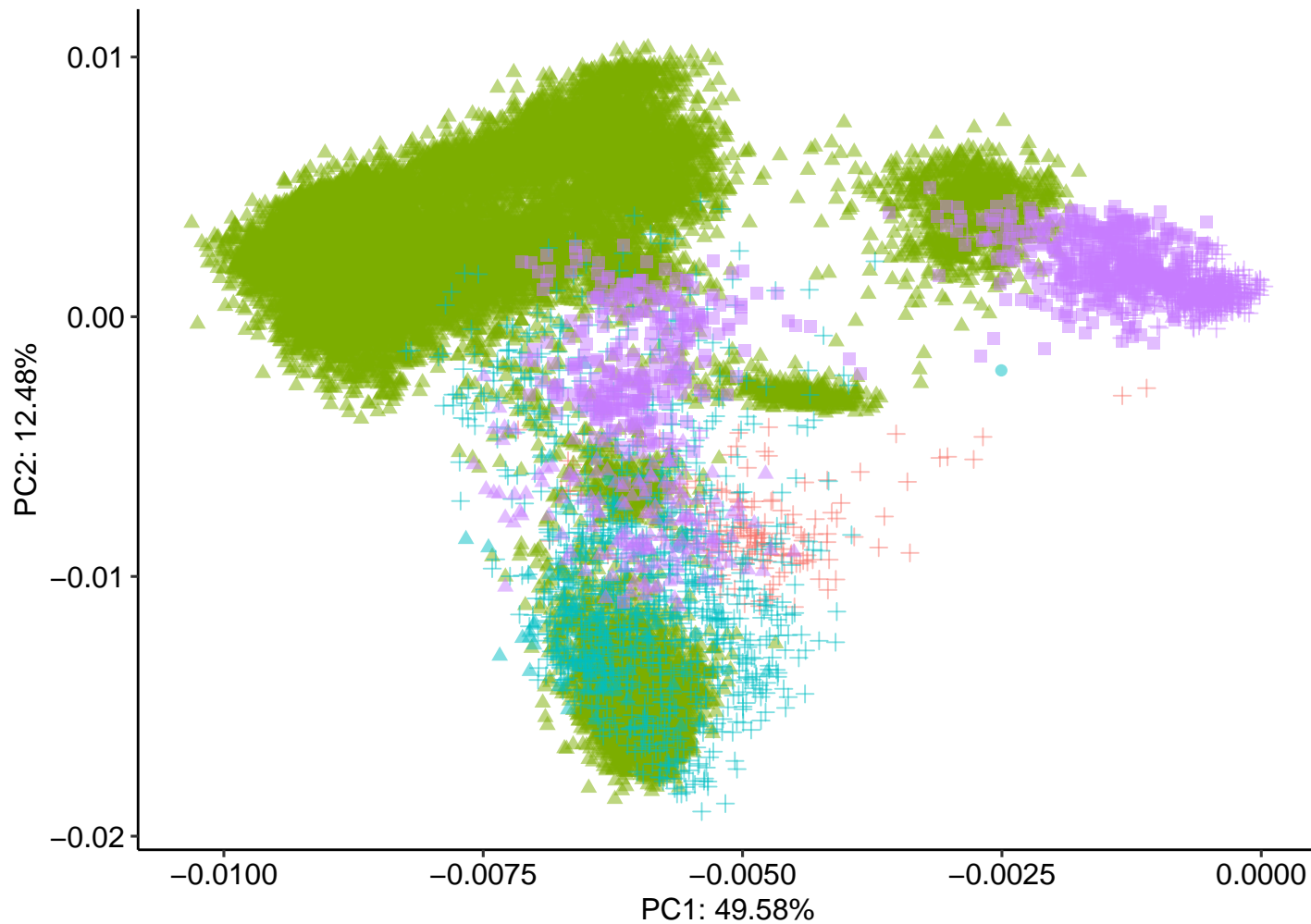
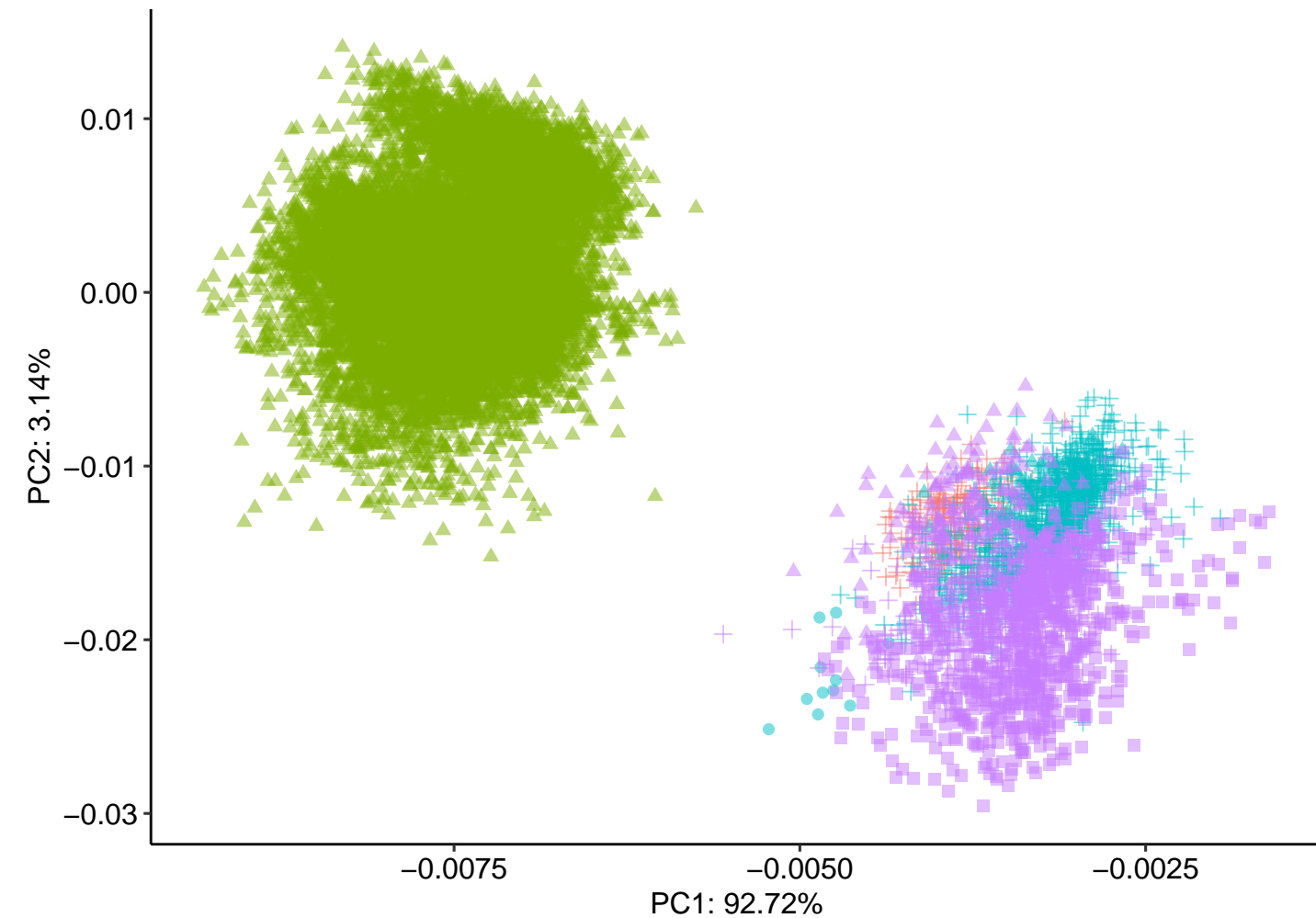


Tumors + Normals: Top variable genes (> 90% quantile) (n = 3666)



Tumors + Normals: House keeping genes (n = 25)



RNA_library ● exome_capture ▲ poly-A ■ poly-A stranded + stranded NA cohort ● GMKF ● GTEx ● PBTA ● TARGET