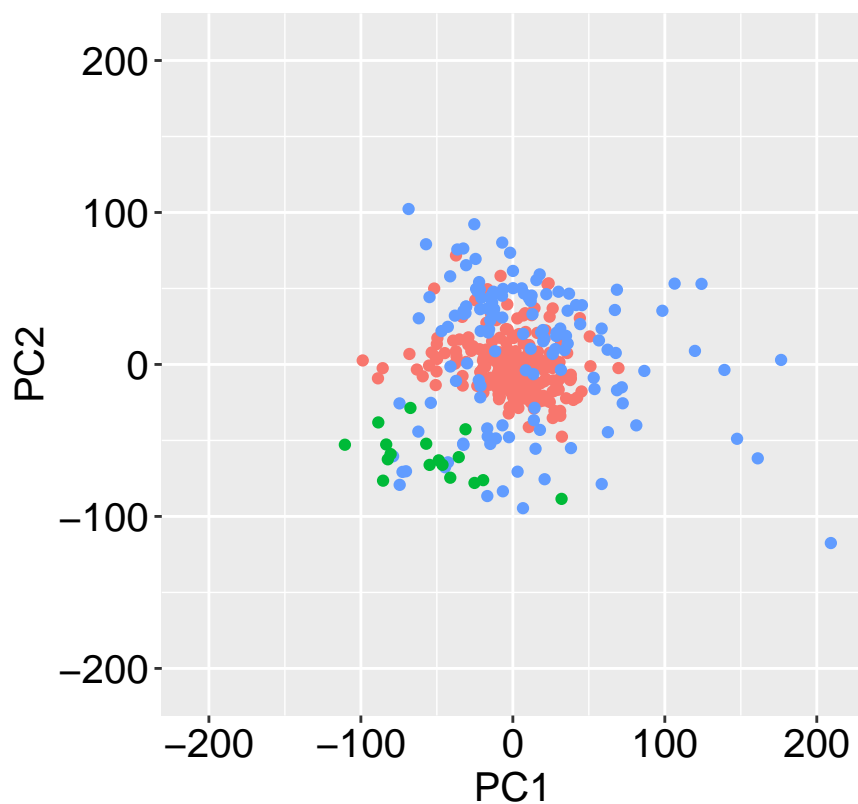


Males



as.factor(sample_type)

- GTEx
- TCGA_normal
- TCGA_tumour