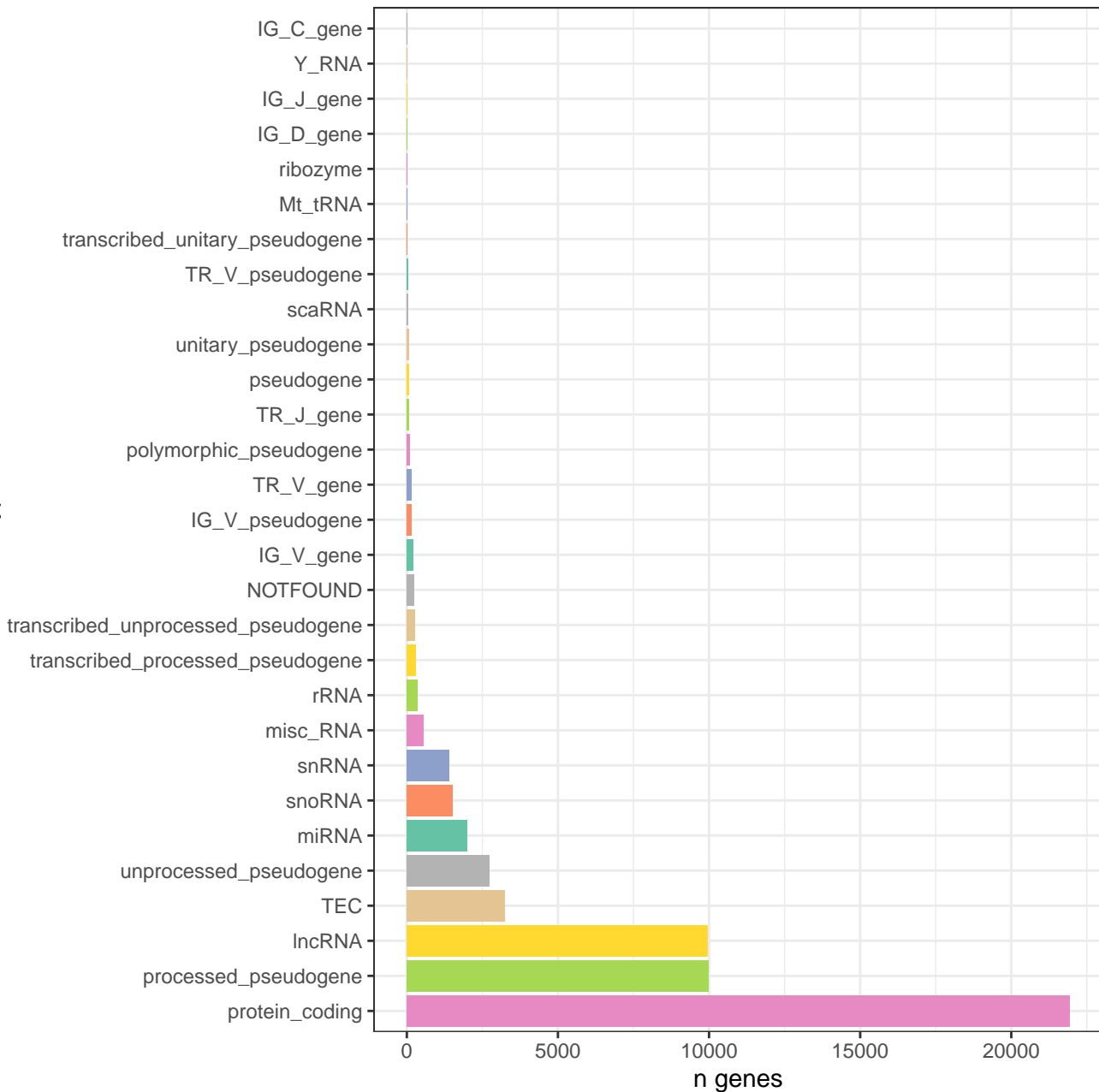
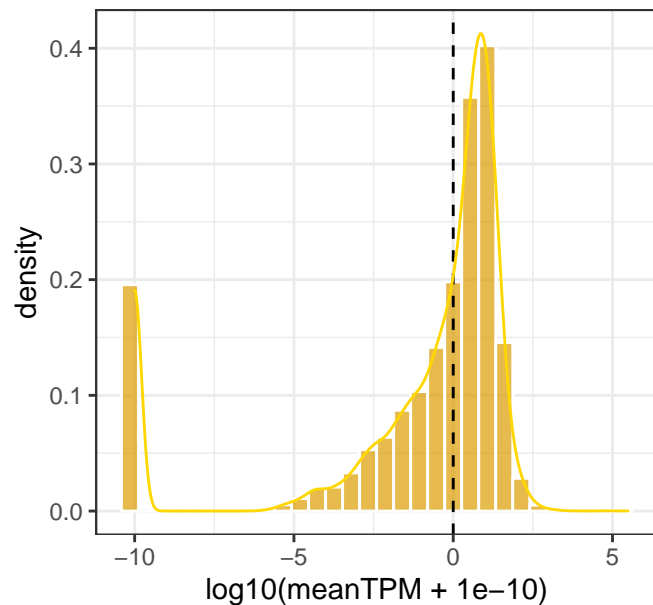


gene biotypes (n genes >10)

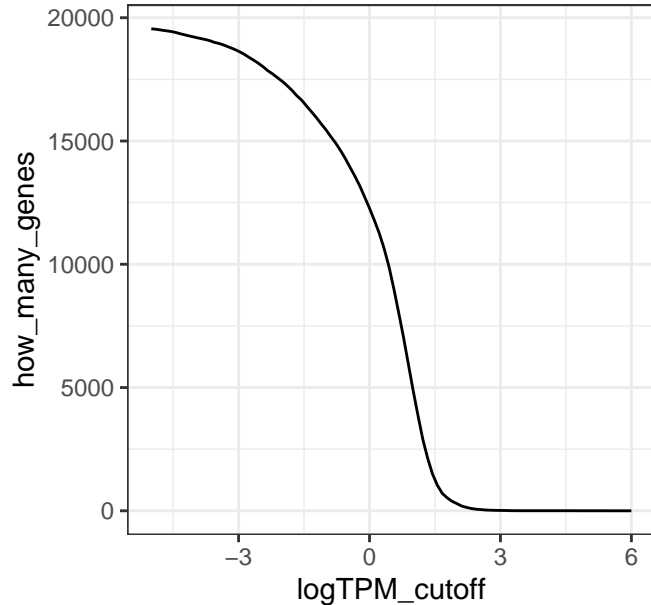
biotype



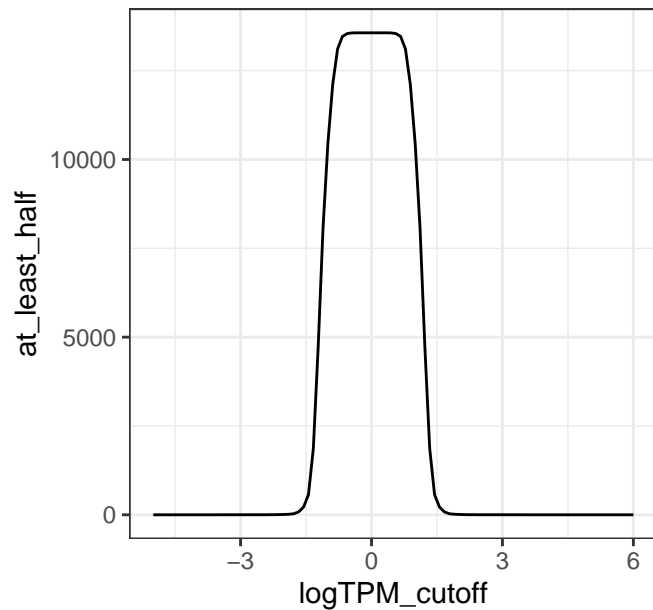
mean TPM distribution across genes



logTPM threshold vs genes preserved



at least one half rule



Raw counts (only protein coding)

