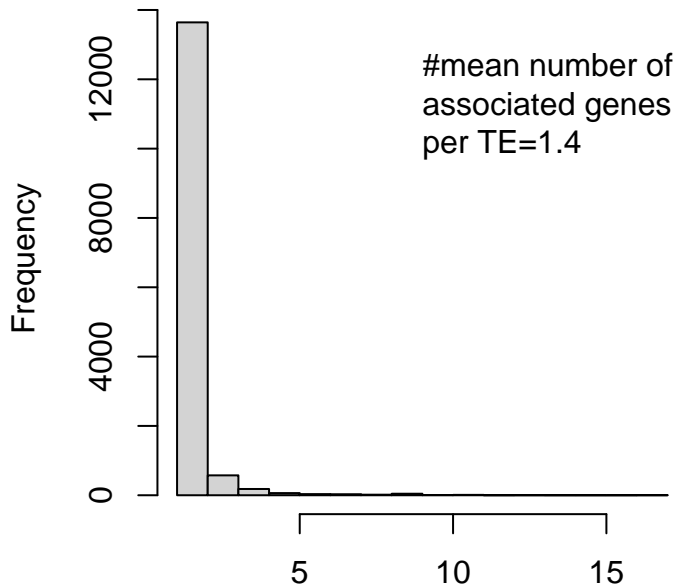
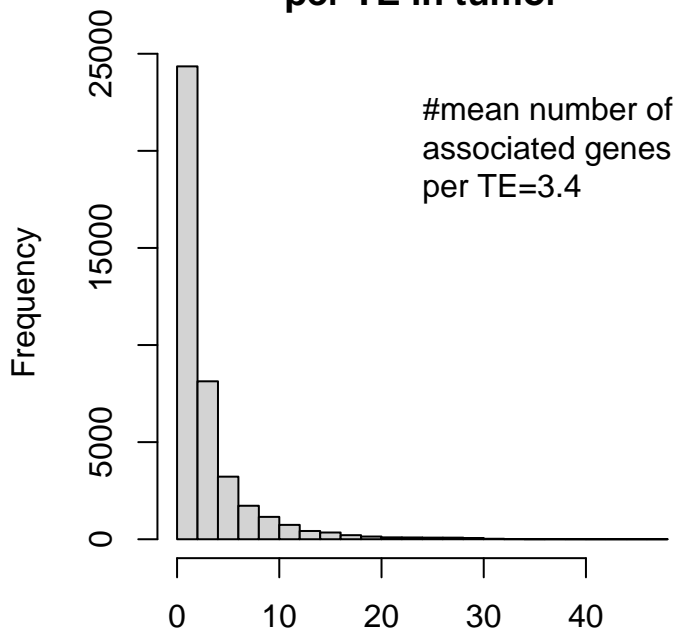


**Number of genes associated
per TE in normal**



Number of genes associated per TE

**Number of genes associated
per TE in tumor**



Number of genes associated per TE