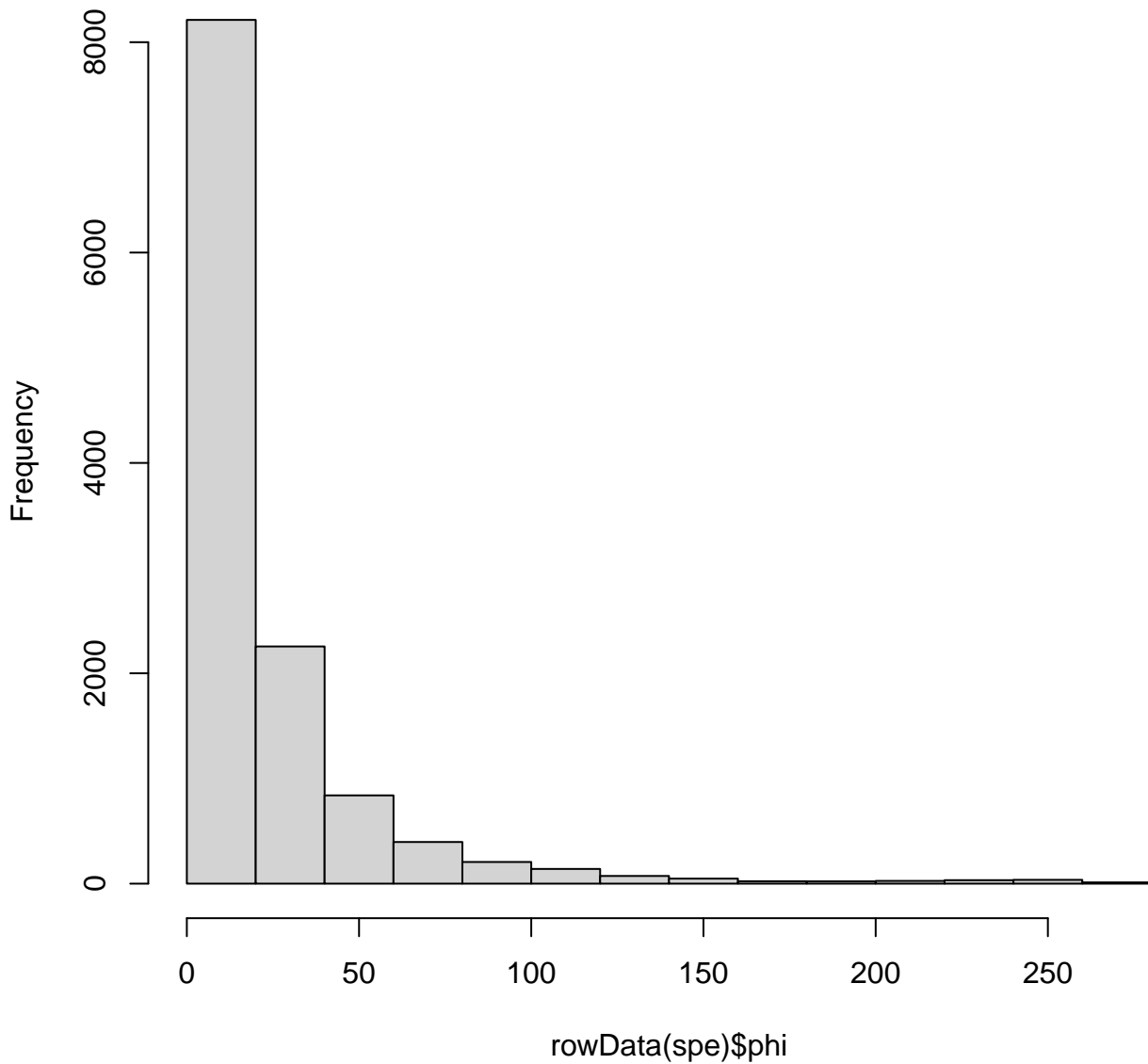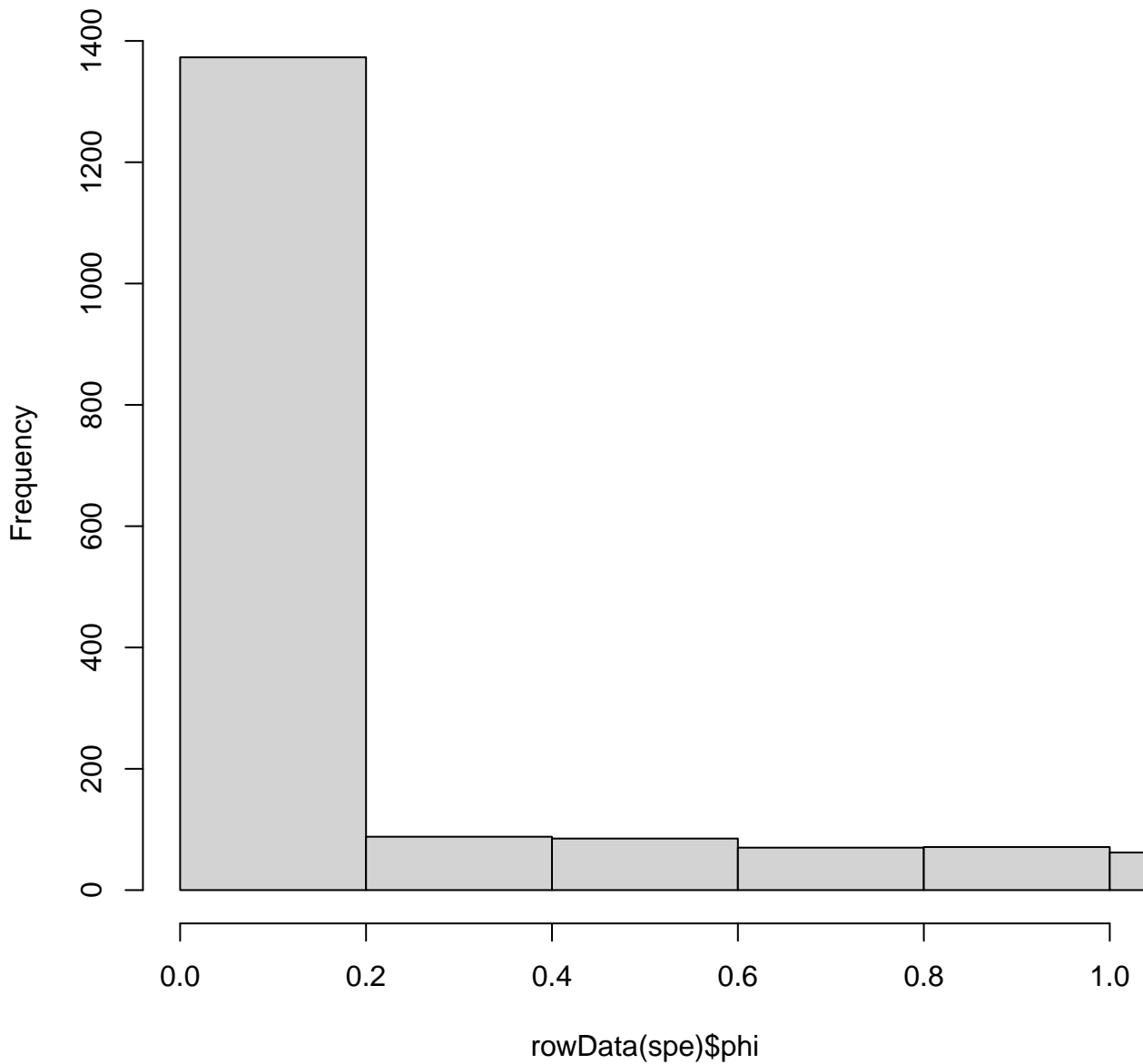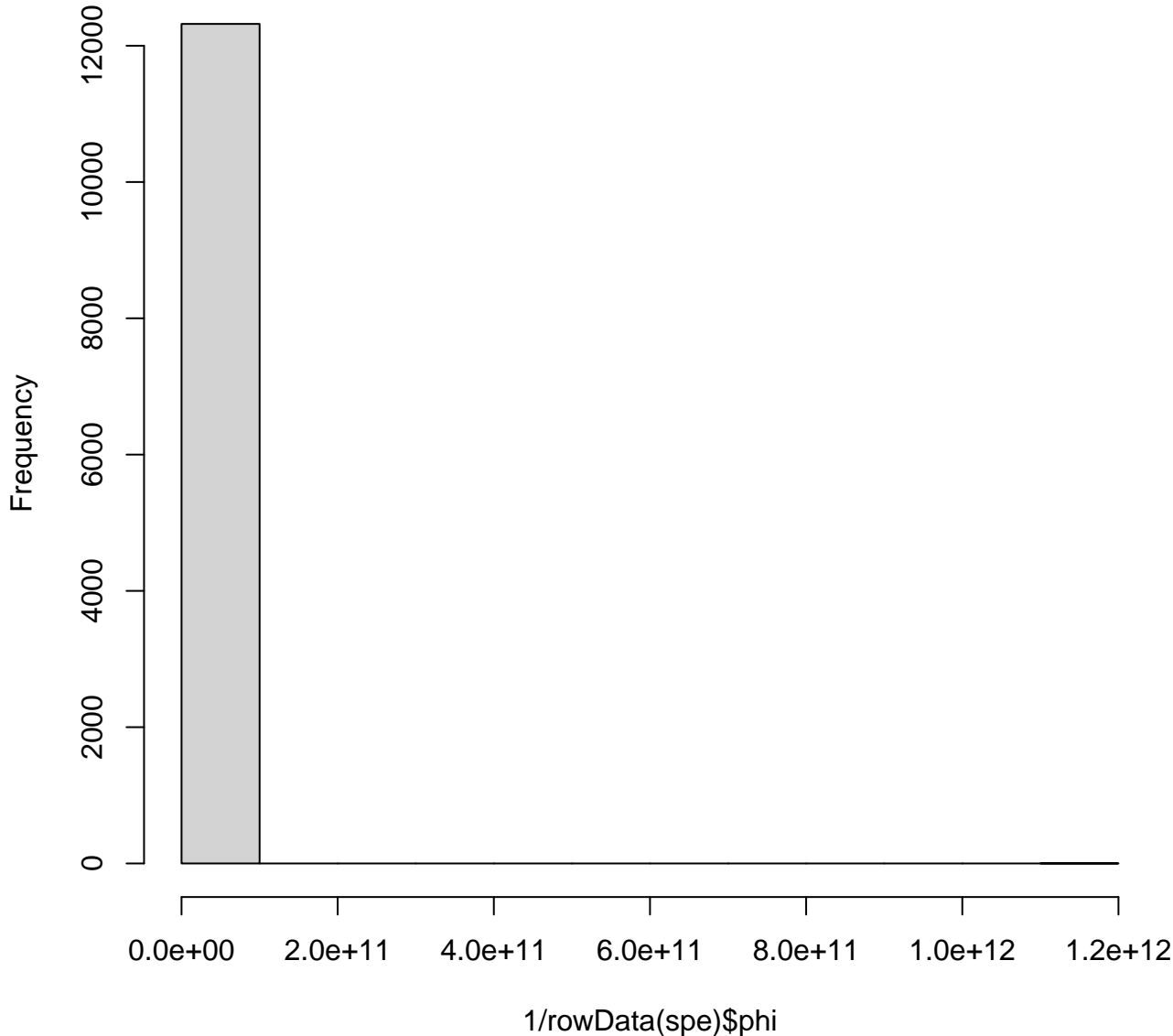**distribution of phi**

distribution of phi

distribution of lengthscale

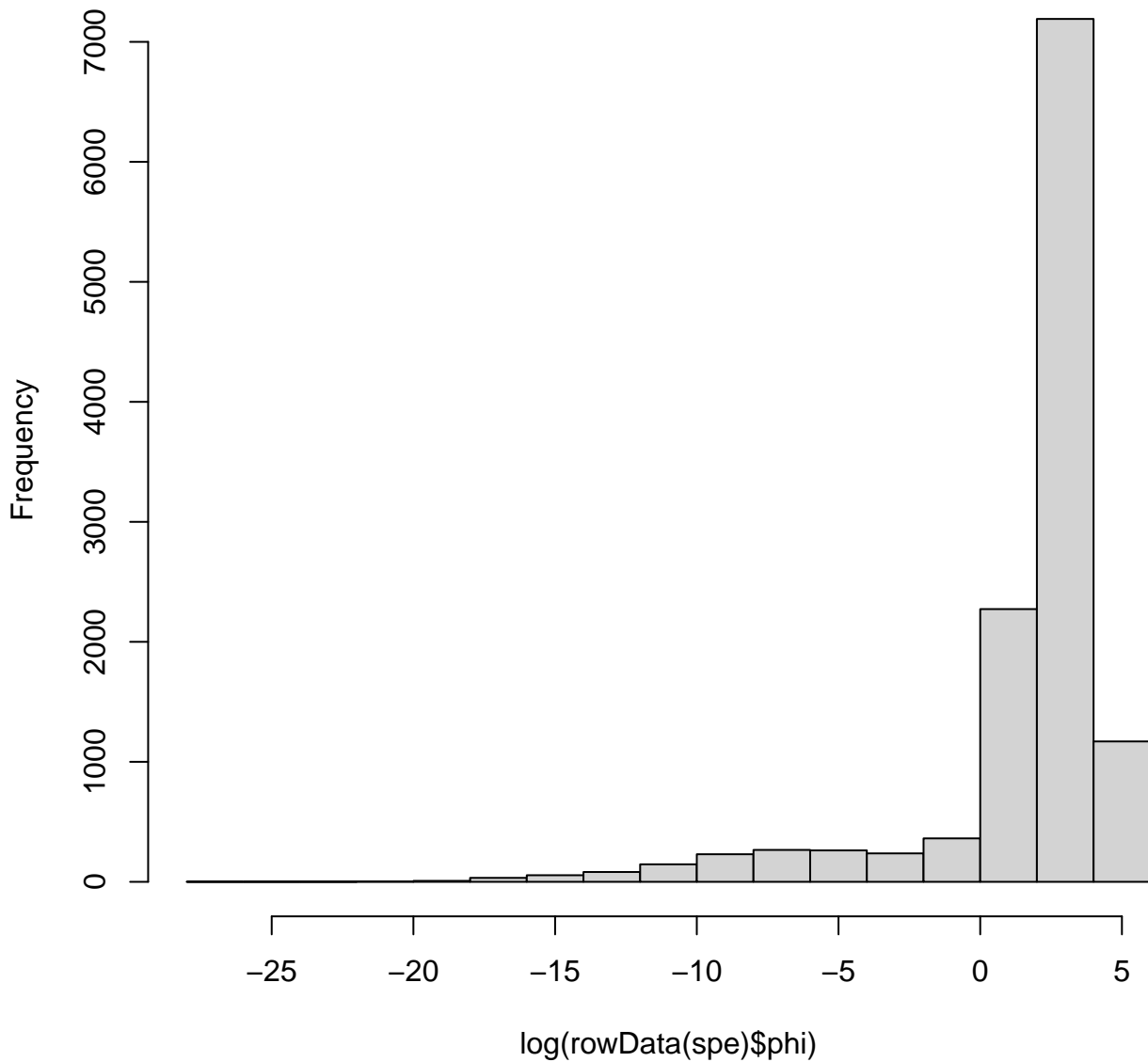**distribution of lengthscale**

Frequency

my_list
removed ~1000 genes with l>10

distribution of lengthscale

Frequency

my_list
removed ~1000 genes with l>10
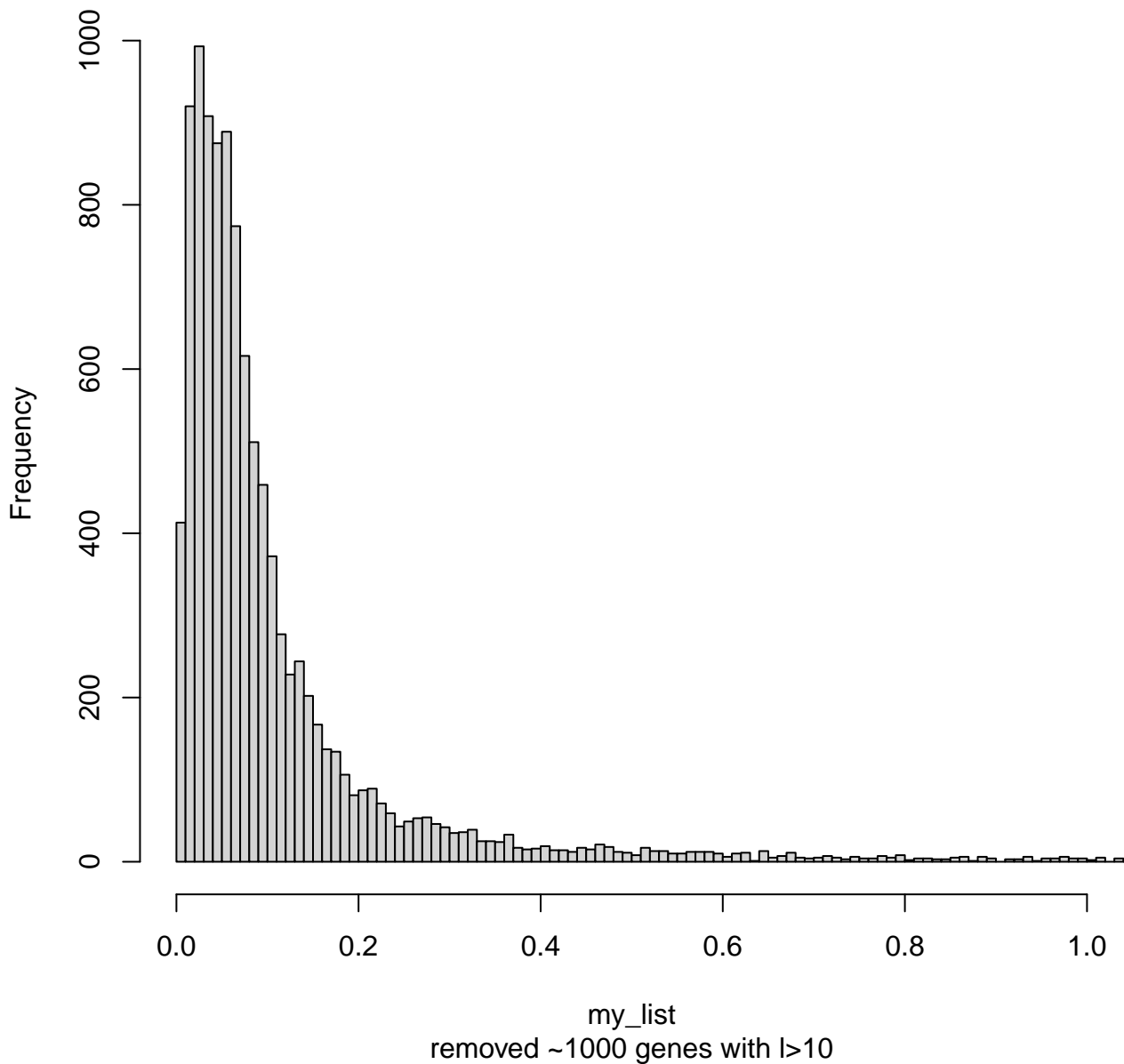
distribution of phi on logscale

log(rowData(spe)$phi)

distribution of lengthscale on logscale