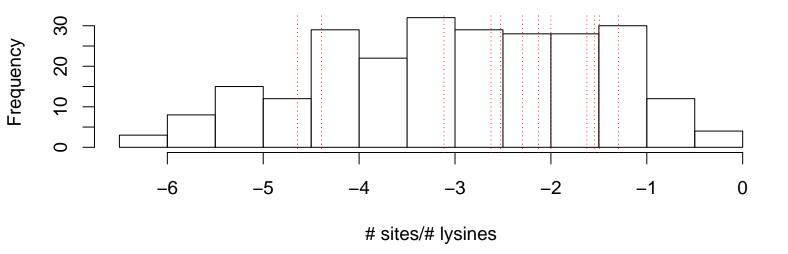
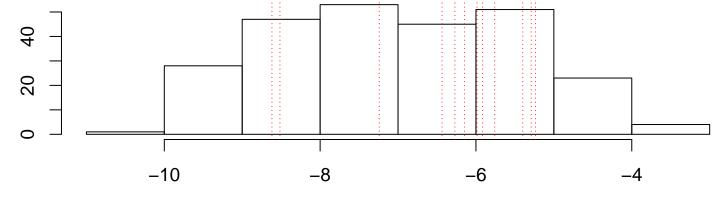
## normalized to number of lysines



## normalized to protein length

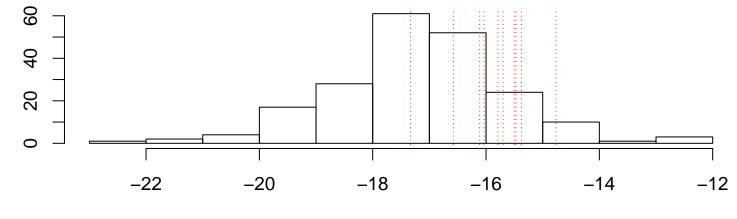


Frequency

Frequency

# sites/protein length

## normalized to protein quant



# sites/ protein quant