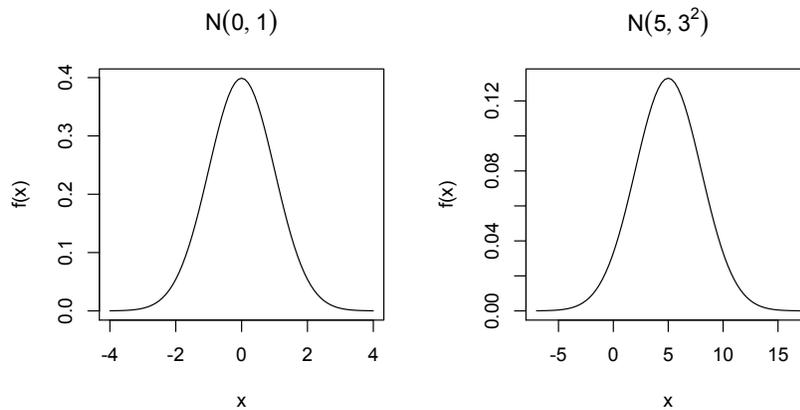


STA 250: STATISTICS  
Lab 4

This lab overviews graphical and numerical summarization of pdfs and pmfs of scalar variables.

**Plotting a pdf or a pmf over an appropriate range**

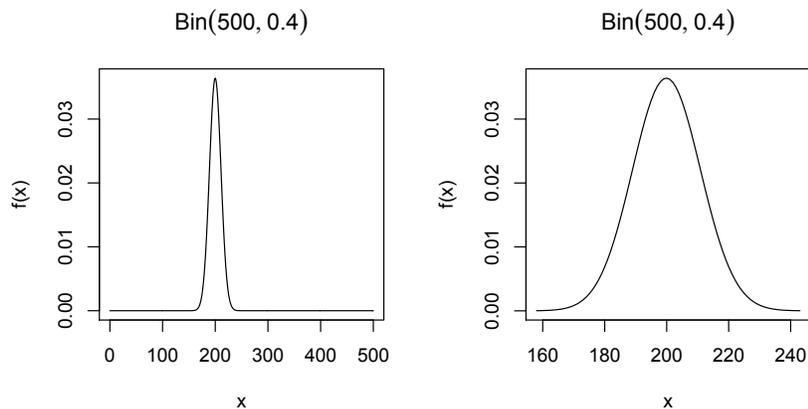
This is a trivial task if you have access to a R function `f()` that returns the pdf/pmf value `f(x)` at any input `x`. The only concern is to choose an appropriate range for `x` to display the curve. To display a `Normal(0, 1)` pdf we could use a range `[-4, 4]`, because most of the area under the `Normal(0, 1)` bell curve is within this range (more than 0.9999). For an arbitrary `Normal( $\mu, \sigma^2$ )` pdf, we can display the range  $\mu \mp 4\sigma$ , i.e., mean plus minus 4 standard deviations, which contains the same area under the `Normal( $\mu, \sigma^2$ )` curve as does `[-4, 4]` for the `Normal(0, 1)` bell curve:



TASK 1. Make a *nice* plot of the `Normal(-500, 402)` pdf. To annotate the axes and add a plot title, use:  
`plot(..., xlab = "x", ylab = "f(x)", main = expression(N(-500, 40^2)))`.

The same strategy applies while plotting other pdfs or pmfs. The `Binomial(500, 0.4)` pmf is supported on the points `0, 1, ..., 500`. But it tightly concentrates around its mean  $500 \times 0.4 = 200$ , and so plotting it over the whole range is rather useless (see the left panel of the Figure below). Instead, we can find a range `[a, b]`, such that pmf puts only a small probability, say  $10^{-4}$ , outside this range. We can find the two end points so that the left out probability is equally split at the two tails. Then we must have `a` as the  $10^{-4}/2$ -th quantile and `b` as the  $(1 - 10^{-4}/2)$ -th quantile of the `Binomial(500, 0.4)` distribution. The following code generates the right panel of the Figure below.

```
a <- qbinom(1e-4 / 2, 500, 0.4)
b <- qbinom(1e-4 / 2, 500, 0.4, lower = FALSE)
x <- a:b
plot(x, dbinom(x, 500, 0.4), ty = "l", ann = FALSE)
title(xlab = "x", ylab = "f(x)", main = expression(Bin(500, 0.4)))
```



TASK 2. What's the role of "lower = FALSE" in the second `qbinom()` call above?

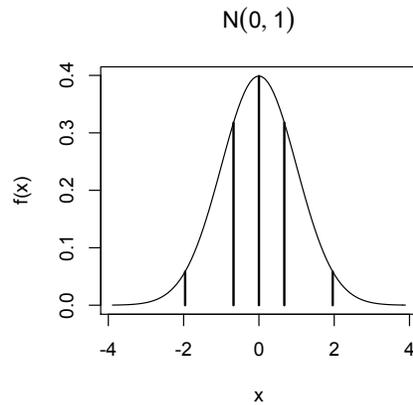
TASK 3. Make plots for pdf/pmf of the `Poisson(30)`, `Gamma(2, 1/5)`, `Logis(4, 10)` and `t4` distributions.

### Quantiles of a distribution

The quantiles of a distribution offer an excellent way to summarize the distribution numerically or graphically. We already saw an example of this above in choosing the range of  $x$  values  $[a, b]$  on which we plot the pdf/pmf. While those give a sense of the extreme, we can also use quantiles to mark the central parts of the distribution.

Recall that for any  $u \in (0, 1)$ , the  $u$ -th quantile of a probability distribution with cdf  $F(x)$  is defined to be the point  $x_u$  such that  $F(x_u) = u$ . That is a vertical slice at  $x = x_u$  splits the total pdf area/pmf total of 1 into  $u$  on the left and  $1 - u$  on the right. The 0.5-th quantile is called the median of the distribution, it splits the pdf area into half and half. The 0.25-th and the 0.75-th quantiles are called the first and the third quartiles (median being the second quartile). Between the 0.025-th and the 0.975-th quantile, the pdf/pmf packs 95% of its area. These five quantiles together give an excellent annotation of a pdf/pmf. The following code and plot show these for `Normal(0, 1)`.

```
qts <- qnorm(c(.025, .25, .5, .75, .975), 0, 1)
a <- qnorm(1e-4/2, 0, 1)
b <- qnorm(1e-4/2, 0, 1, lower = FALSE)
x <- seq(a, b, length = 101)
plot(x, dnorm(x, 0, 1), ty = "l", ann = FALSE)
title(xlab = "x", ylab = "f(x)", main = expression(N(0,1)))
segments(qts, 0 * qts, qts, dnorm(qts, 0, 1), lwd = 2)
```



TASK 4. Repeat task 3 now with the five-quantile summaries overlaid on each pdf/pmf.

### Getting quantiles from samples

When R does not provide a quantile function for a distribution, we can still approximate its quantiles by drawing random samples from it. The code below compares a direct calculation against a sample based approximation of quantiles for  $\text{Normal}(0, 1)$

```
> qnorm(c(.025, .25, .5, .75, .975), 0, 1)
[1] -1.96 -0.67 0.00 0.67 1.96
> as.numeric(quantile(rnorm(1e4), c(.025, .25, .5, .75, .975)))
[1] -1.995 -0.690 -0.015 0.656 1.900
```

TASK 5. Get five-quantile summaries of  $\text{Poisson}(30)$ ,  $\text{Gamma}(2, 1/5)$ ,  $\text{Logis}(4, 10)$  and  $t_4$  based on random samples drawn from these distributions (use  $10^4$  draws) and compare them with the actual values.

Consider the model  $X \sim \text{Binomial}(20, p)$  with prior  $\xi(p) = \text{Beta}(10, 10)$ . In lecture we discussed that this defines a joint pd/mf  $h(x, p)$  for  $(X, p)$ . From the joint we could extract the marginal pmf for  $X$ . While it is possible to write down a formula for this marginal pmf, it is much easier to draw samples of  $X$  from this marginal pmf through the following:

```
p <- rbeta(1, 10, 10)
x <- rbinom(1, 20, p) ## using the 'p' generated above
```

which you can wrap inside `replicate()` to generate many samples at one go.

TASK 6. Use the above code with `replicate()` to generate  $10^4$  samples of  $X$  from its marginal and get their five-quantile summary.