1. Write a pois.prob() function that computes P(X=x), $P(X \neq x)$, P(X < x), $P(X \leq x)$, P(X > x), and $P(X \geq x)$. Enable the user to specify the rate parameter λ .

Solution: Below is the code for computing various probabilities for the Poisson Distribution. The **prob** parameter was removed as a parameter of the function, as it would only be used in calculations involving **qpois**, the inverse CDF, which was not used in the function. λ was passed in directly as a parameter for the Poisson Distribution, and statements were added to make sure λ was ≥ 0 and x was in the support.

```
pois.prob <- function(x, lambda, type="<="){</pre>
  if (lambda < 0){</pre>
    # lambda must be <= 0
    lambda = 0
  if (x < 0)
    # x must be <= 0
    x = 0
  if (type == "="){
    # definition of PMF
    return(dpois(x, lambda))
  if (type == "!="){
    # complement rule
    return(1-dpois(x, lamda))
  if(type == "<="){
    # definition of CDF
    return(ppois(x, lambda))
  if (type == "<"){
    \# P(X < x) = P(X \le x-1)
    return(ppois(x-1, lambda))
  if (type == ">="){
# P(X >= x) = 1 - P(X <= x-1)
    return(1-ppois(x-1, lambda))
  if (type == ">") {
# P(X > x) = 1 - P(X <= x)
    return(1-ppois(x, lambda))
```

2. Write a beta.prob() function that computes P(X=x), $P(X\neq x)$, P(X< x), $P(X\leq x)$, P(X>x), and $P(X\geq x)$ for a beta distribution. Enable the user to specify the shape parameters α and β .

```
beta.prob <- function(x, size, prob, type="<="){
    # Use dbeta and pbeta to conditionally return the correct probability
}</pre>
```