

# bayesian\_stats\_ch2

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```
# help(rpois)

df1 <- tibble(x = rpois(n = 100, lambda = 2.1),
              y = rpois(n = 100, lambda = 21),
              label = "n = 100")

df2 <- tibble(x = rpois(n = 1000, lambda = 2.1),
              y = rpois(n = 1000, lambda = 21),
              label = "n = 1000")

df3 <- tibble(x = rpois(n = 10000, lambda = 2.1),
              y = rpois(n = 10000, lambda = 21),
              label = "n = 10000")

df <- df1 %>%
  bind_rows(df2) %>%
  bind_rows(df3)

ggx <- df %>%
  ggplot(aes(x = x, fill = factor(label)))

ggy <- df %>%
  ggplot(aes(x = y, fill = factor(label)))

gg_x_pois <- ggx +
  geom_histogram() +
  facet_wrap(~factor(label)) +
  labs(title = "Poisson(lambda = 2.1)")
```

```

gg_y_pois <- ggy +
  geom_histogram() +
  facet_wrap(~factor(label)) +
  labs(title = "Poisson(lambda = 21)")

p_load(cowplot, ggpubr)

# cowplot::plot_grid(gg_x_pois, gg_y_pois, align = "h")

gg_poisson <- ggpubr::ggarrange(gg_x_pois, gg_y_pois, nrow = 2, ncol = 1)

## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.
## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.

ggsave(filename = "poisson_dist_ch2.png", plot = gg_poisson)

## Saving 6.5 x 4.5 in image

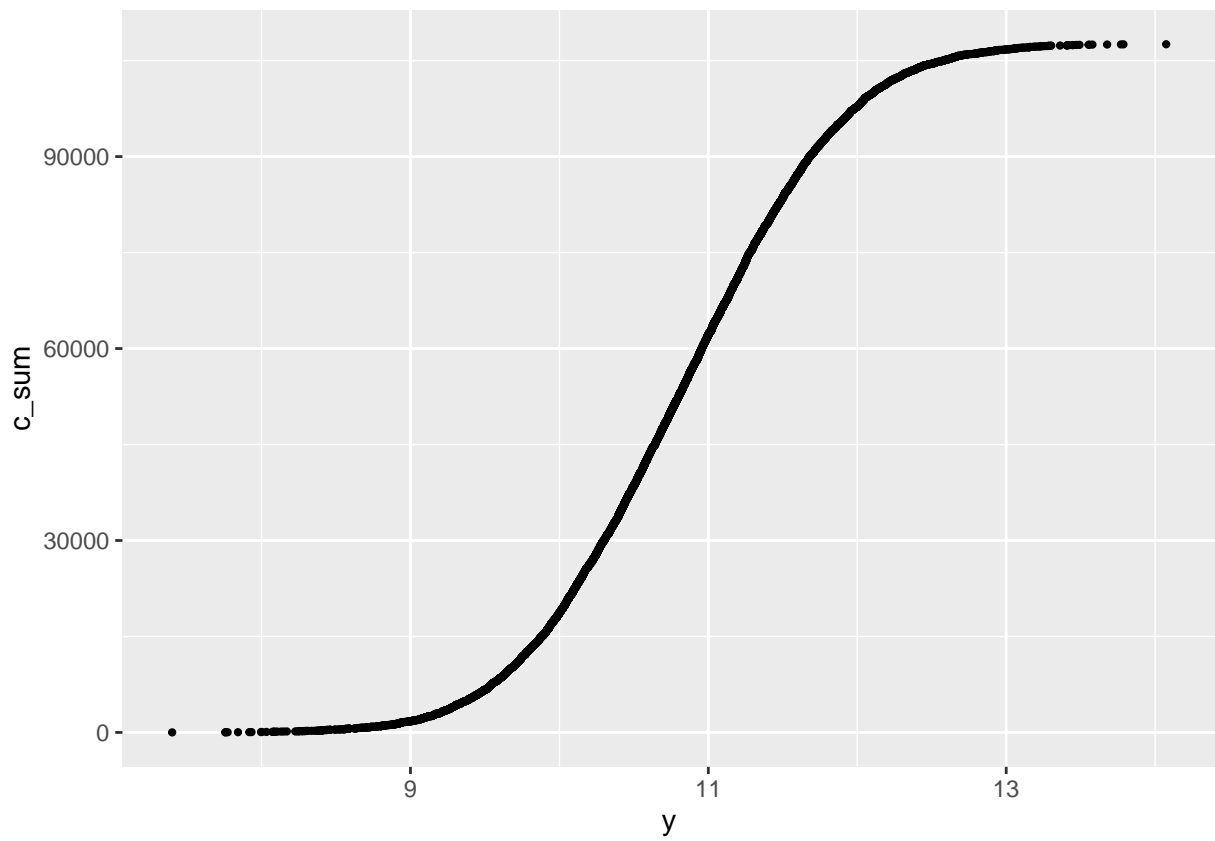
y <- rnorm(n = 10000, mean = 10.75, sd = sqrt(0.8))

empirical_dist <- cumsum(sort(y))

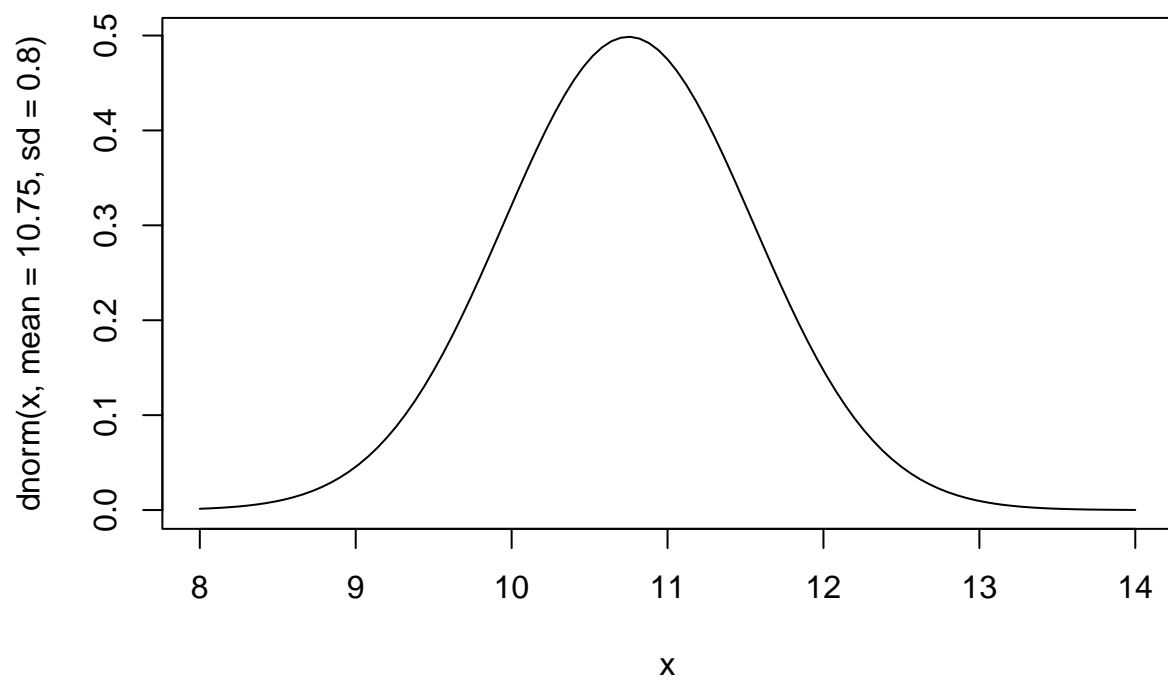
df <- tibble(c_sum = empirical_dist,
             y = sort(y))

df %>%
  ggplot(aes(x = y, y = c_sum)) +
  geom_point(size = 0.8)

```



```
curve(dnorm(x, mean = 10.75, sd = 0.8), from = 8, to = 14)
```



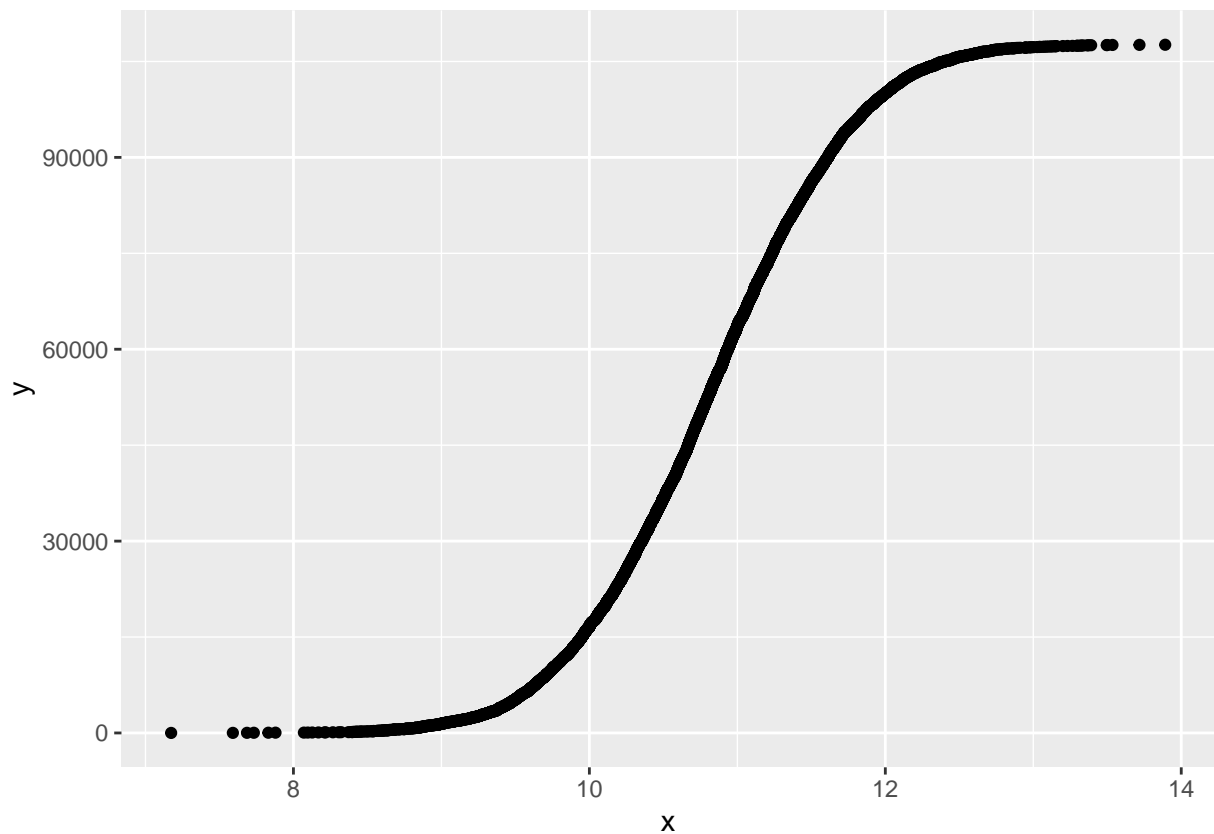
```

cdf <- ecdf(rnorm(n = 10000, mean = 10.75, sd = 0.8))

df <- tibble(x = sort(rnorm(n = 10000, mean = 10.75, sd = 0.8)), y = cumsum(x))

df %>%
  ggplot(aes(x = x, y = y)) +
  geom_point()

```



```

# help(ecdf)
# help(dnorm)

```

```

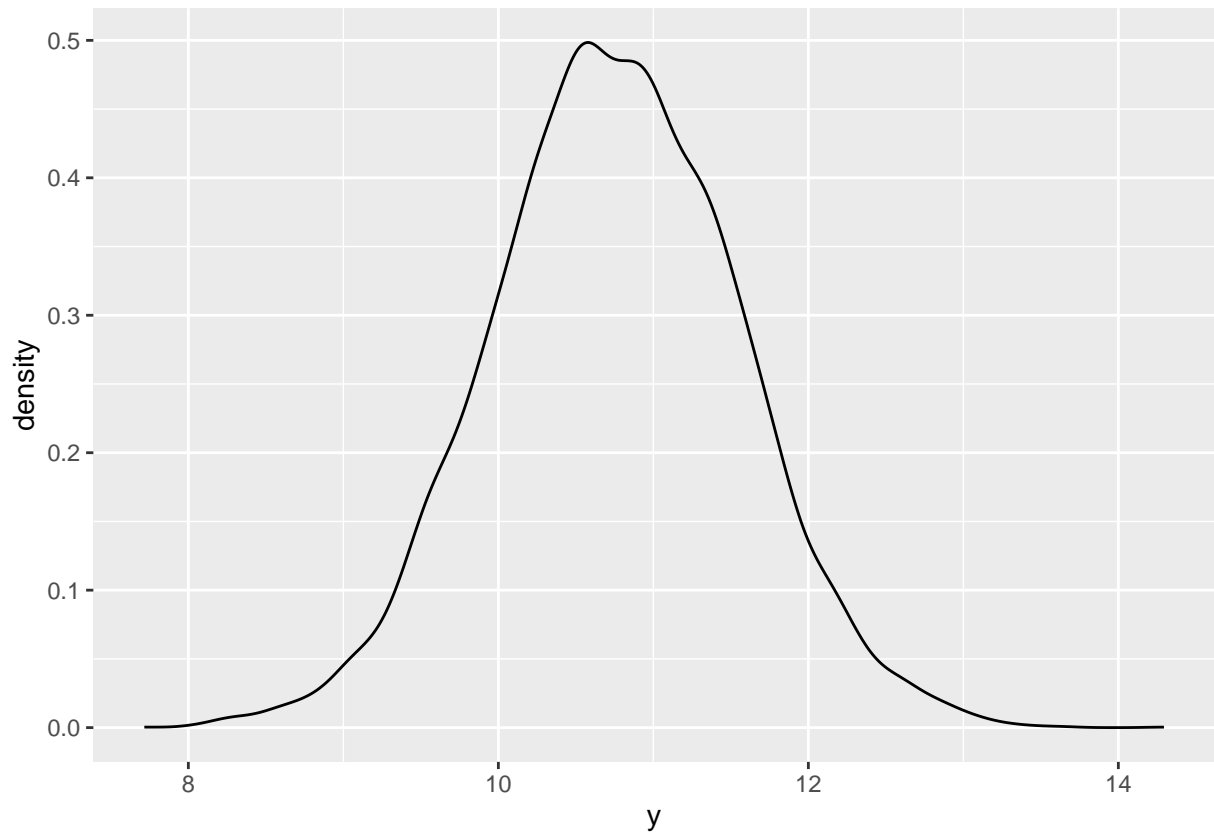
n = 10000

df <- tibble(y = rnorm(n = n, mean = 10.75, sd = 0.8),
             logy = rlnorm(n = n, meanlog = 10.75, sdlog = 0.8),
             d = dnorm(y),
             c = cumsum(d))

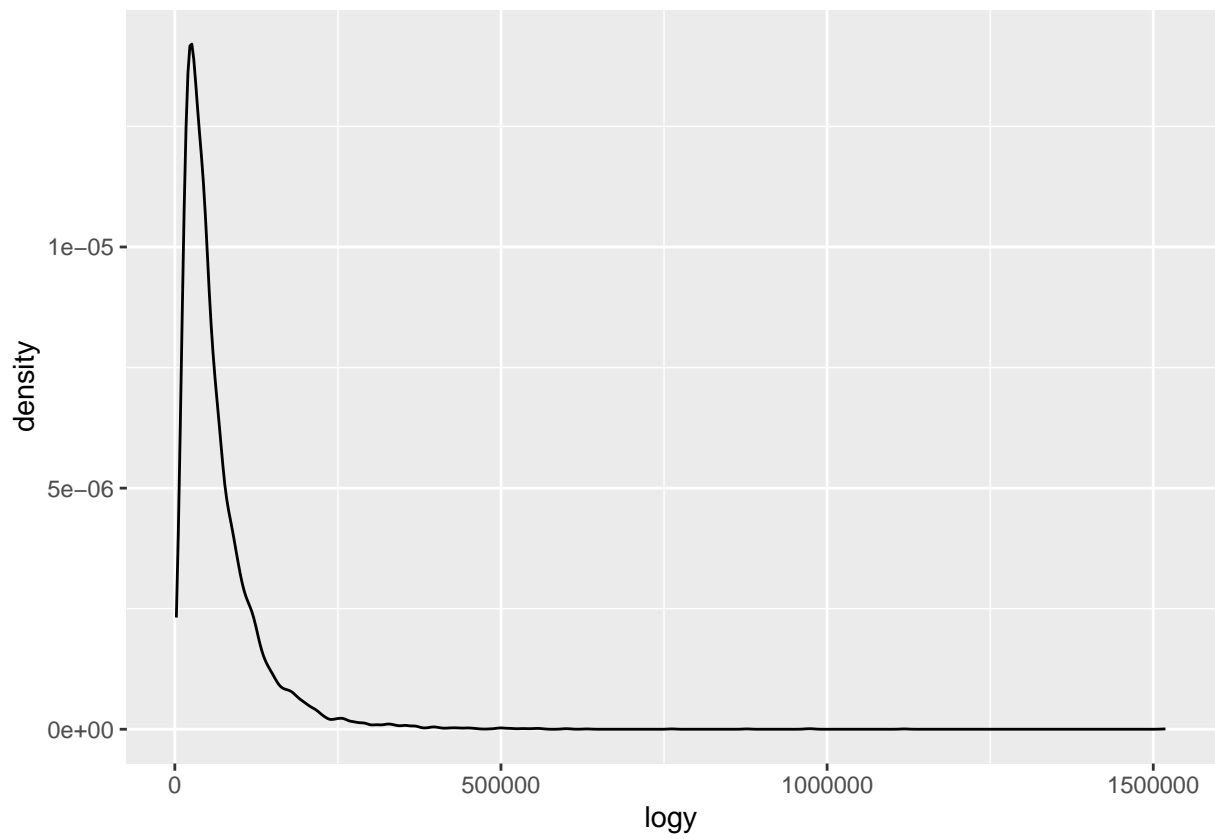
df %>%
  ggplot(aes(x = y)) +

```

```
geom_density()
```



```
df %>%  
  ggplot(aes(x = logy)) +  
  geom_density()
```



```
# help(pnorm)
```