

My M&M OCD

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02 04, 2025

Intro

The goal of this simulation is to test the statistics of M&M and other stacks even Chocolate lentils by color, I wanted to know, if I eat m&m package 2 by 2, separated by color, what is the chance of my finishing the package without mixing any color in one bite.

In addition, here are some BI incite that needed to be checked:

1. What is the probability of M&M packages packaged fairly?
2. What is the probability of M&M packages packaged without one color?
3. How does the size of the package or number of colors affect this probability?

The method is based of simulation of some M&M bags, according to the most common sizes. Each time we sample x lentils, name them by colors (represented as factorial numbers), and see the results for many packages as a statistic data.

Parameters

```
## -- Attaching core tidyverse packages ----- tidyverse 2.0.0 --
## v dplyr      1.1.4      v readr      2.1.5
## v forcats    1.0.0      v stringr    1.5.1
## v ggplot2     3.5.1      v tibble     3.2.1
## v lubridate  1.9.4      v tidyr      1.3.1
## v purrr       1.0.4
## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()     masks stats::lag()
## i Use the conflicted package (<http://conflicted.r-lib.org/>) to force all conflicts to become errors
##
## Attaching package: 'MASS'
##
##
## The following object is masked from 'package:dplyr':
##
##   select
##
##
```

```
## Attaching package: 'scales'
##
##
## The following object is masked from 'package:purrr':
##
##   discard
##
##
## The following object is masked from 'package:readr':
##
##   col_factor
```

The basic parameters (will be changed later):

```
#parameters
nn<- 500           #numbers of bags per sample
n_color<- 6        #unique colors of M&M
gram<- 0.91        #weight of one M&M
bag_g<- 250        #common weight of M&M package
n_unit<- bag_g/gram #M&M per package
av_per_color= n_unit/n_color
paste0("The avarage number of lentils per color is ", round(av_per_color,2))
```

```
## [1] "The avarage number of lentils per color is 45.79"
```

Creating of Sample

General Sample

create_bag is a function to create one snack package as matrix.

sample_MnM is a function to create n bags from the create_bag function.

```
## [1] "One bag:"
```

```
##      1  2  3  4  5  6
## [1,] 11 21 15 15 16 23
```

```
## [1] "3 bags:"
```

```
##      1 2 3 4 5 6
## Bag_1 2 2 0 2 2 2
## Bag_2 4 3 0 0 1 3
## Bag_3 0 2 1 3 2 2
```

Preview Graph

Now will be creating nn bugs of M&M

```

MnM_sample<- sample_MnM(nn,n_unit,n_color)

MnM_sample<-
  MnM_sample %>% as_data_frame() %>%
  mutate(even_count= rowSums(across(everything() , ~ .x %% 2 == 0)), #how many evens colors there are
         even_evens= rowSums(across(c(1:6), ~ .x %% 2 == 1)) %% 2 ==0, #are the uneven colors even
         Variance=      apply(across(c(1:6)), 1, var), #var of candy per color/ type
         low_col=      rowSums(across(c(1:6), ~ .x <= 0.6*av_per_color)),
         min=          apply(across(c(1:6)),1,FUN = min)) # lowest value in color

MnM_sample %>% head(6)

```

```

## # A tibble: 6 x 11
##       V1     V2     V3     V4     V5     V6 even_count even_evens Variance low_col
##   <int> <int> <int> <int> <int> <int>    <dbl> <lgl>      <dbl>    <dbl>
## 1    25    48    50    59    48    45         3 FALSE      127.         1
## 2    53    49    42    53    38    40         3 FALSE      44.6         0
## 3    45    46    45    45    45    48         2 TRUE       1.47         0
## 4    57    48    39    45    44    42         3 FALSE      39.0         0
## 5    43    44    51    48    43    45         2 TRUE       10.3         0
## 6    51    41    49    49    40    44         2 TRUE       21.5         0
## # i 1 more variable: min <int>

```

ggplot the MnM sample sample

```

MnM_sample$even_count %>%
table()

```

```

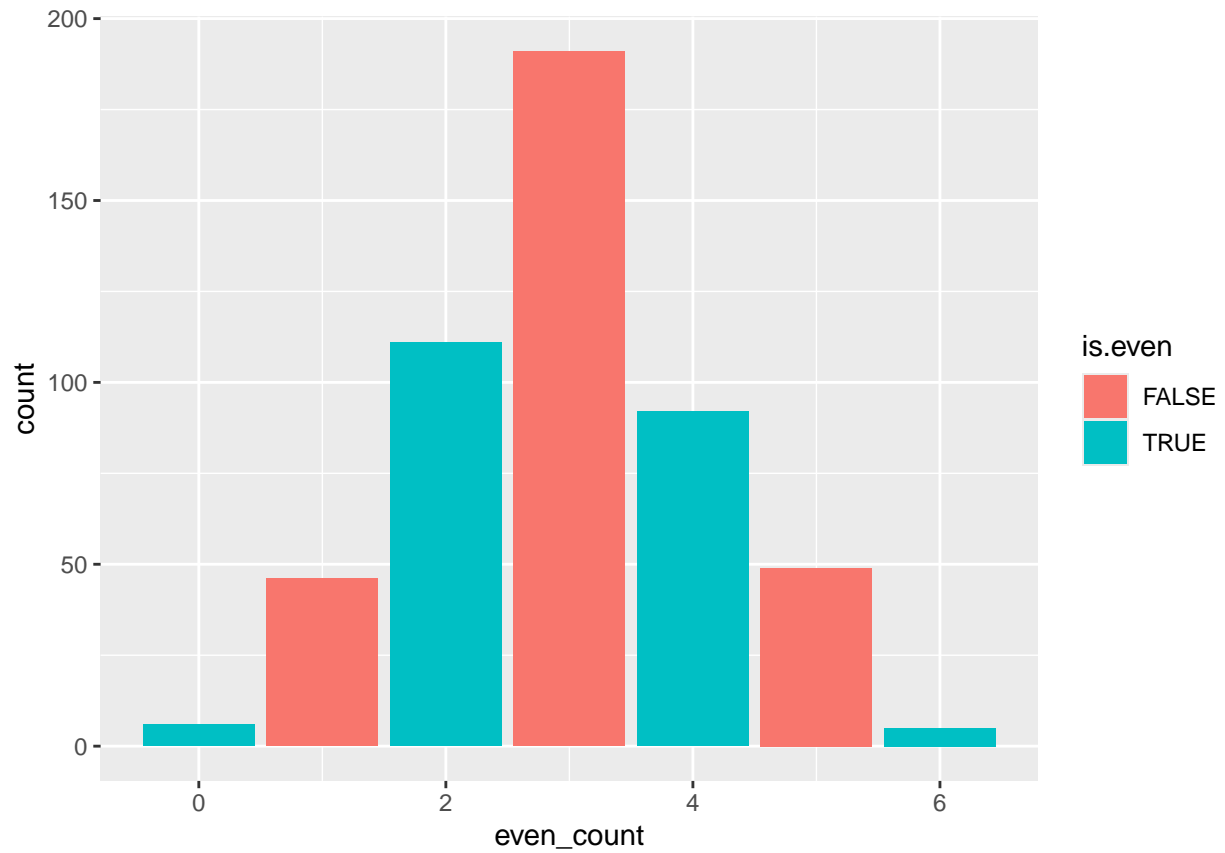
## .
##  0  1  2  3  4  5  6
##  6 46 111 191 92 49 5

```

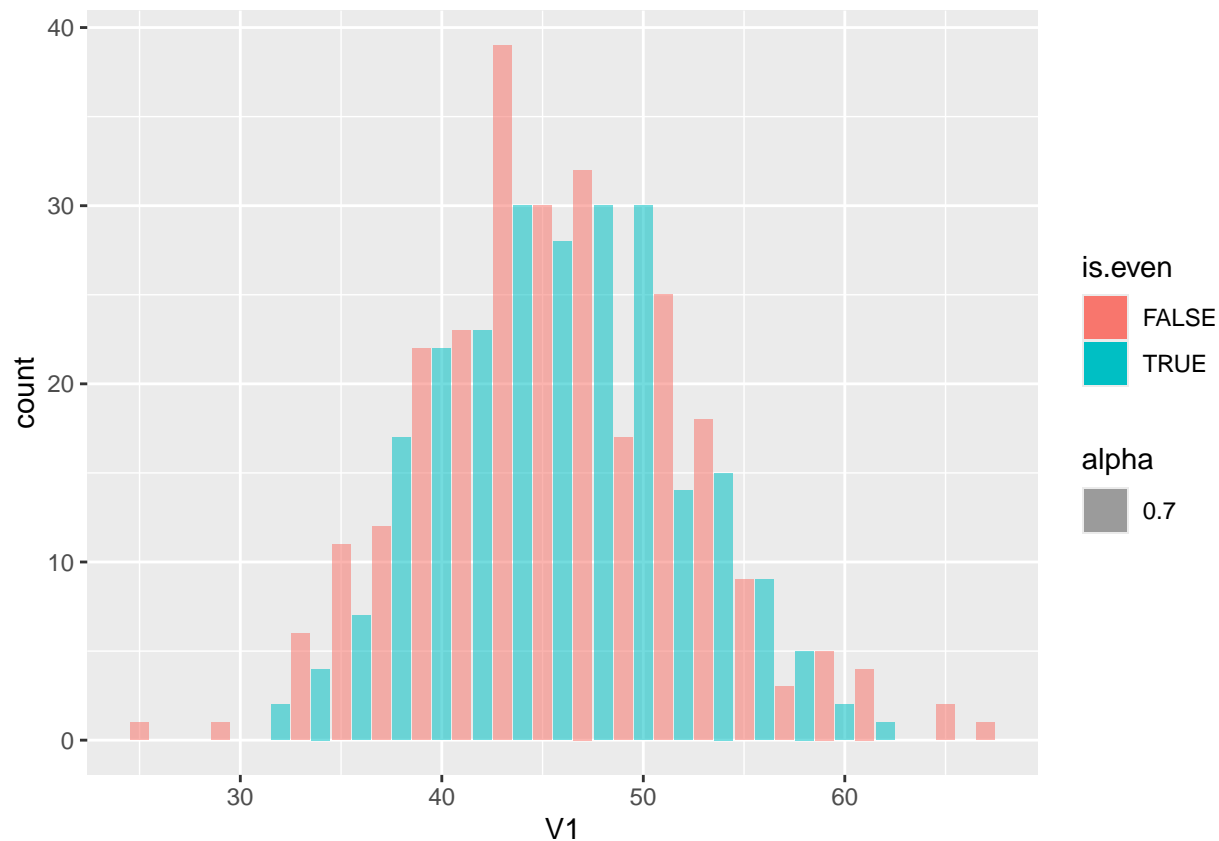
```

MnM_sample %>%
  mutate(is.even = even_count %%2 ==0) %>%
  ggplot(aes(x= even_count, fill= is.even))+
  geom_bar()

```



```
MnM_sample %>%  
  mutate(is.even = V1 %%2 ==0) %>%  
  ggplot(aes(x= V1, fill= is.even, alpha= 0.7))+  
  geom_bar()
```



```
#summary of all colors
rbind(
MnM_sample$V1 %>% summary(),
MnM_sample$V2 %>% summary(),
MnM_sample$V3 %>% summary(),
MnM_sample$V4 %>% summary(),
MnM_sample$V5 %>% summary(),
MnM_sample$V6 %>% summary()
) %>% as.data.frame() %>% cbind(sapply(MnM_sample[,1:6],var,na.rm=1)) %>%
  rename("Var" ="sapply(MnM_sample[, 1:6], var, na.rm = 1)")
```

##	Min.	1st Qu.	Median	Mean	3rd Qu.	Max.	Var
## V1	25	41	45.5	45.766	50	67	40.30786
## V2	31	42	46.0	46.026	50	65	37.78890
## V3	30	42	46.0	46.004	50	68	38.38075
## V4	27	41	45.0	45.276	50	68	43.12608
## V5	30	42	46.0	45.766	50	61	35.76277
## V6	29	41	46.0	45.734	50	64	39.67860

Test Expected Value

to see if the μ of the lentils per color are fair, we will test it per column with t.test.

```

#test mu is av_per_color
check_mean_hypothesis <- function(data, column_name, X) {
  test_result <- t.test(data[[column_name]], mu = X)
  return(test_result$p.value)
}

columns_to_test <- colnames(MnM_sample[,1:6])
test_results <- sapply(columns_to_test, function(col) {
  check_mean_hypothesis(MnM_sample, col, av_per_color)
})

t(t(test_results))

```

```

##           [,1]
## V1 0.93954148
## V2 0.38615243
## V3 0.43502140
## V4 0.08215875
## V5 0.93582236
## V6 0.84932521

```

```

check_mean_hypothesis <- function(data, column_name, X) {
  t.test(data[[column_name]], mu = X)$p.value}

check_two_sample_t_test <- function(data, col1, col2) {
  t.test(data[[col1]], data[[col2]])$p.value}

columns_to_test <- colnames(MnM_sample[,1:6])
num_cols <- length(columns_to_test)
p_value_matrix <- matrix(NA, nrow = num_cols, ncol = num_cols, dimnames = list(columns_to_test, columns_to_test))
for (i in 1:num_cols) {
  for (j in 1:num_cols) {
    if (i == j) { #diagonal test of mu
      p_value_matrix[i, j] <- check_mean_hypothesis(MnM_sample, columns_to_test[i], av_per_color)
    } else { # 2 sample t.test
      p_value_matrix[i, j] <- check_two_sample_t_test(MnM_sample, columns_to_test[i], columns_to_test[j])
    }
  }
}

round(as.data.frame(p_value_matrix),4)

```

```

##           V1      V2      V3      V4      V5      V6
## V1 0.9395 0.5108 0.5487 0.2306 1.0000 0.9362
## V2 0.5108 0.3862 0.9551 0.0626 0.4980 0.4584
## V3 0.5487 0.9551 0.4350 0.0717 0.5367 0.4946
## V4 0.2306 0.0626 0.0717 0.0822 0.2176 0.2607
## V5 1.0000 0.4980 0.5367 0.2176 0.9358 0.9344
## V6 0.9362 0.4584 0.4946 0.2607 0.9344 0.8493

```

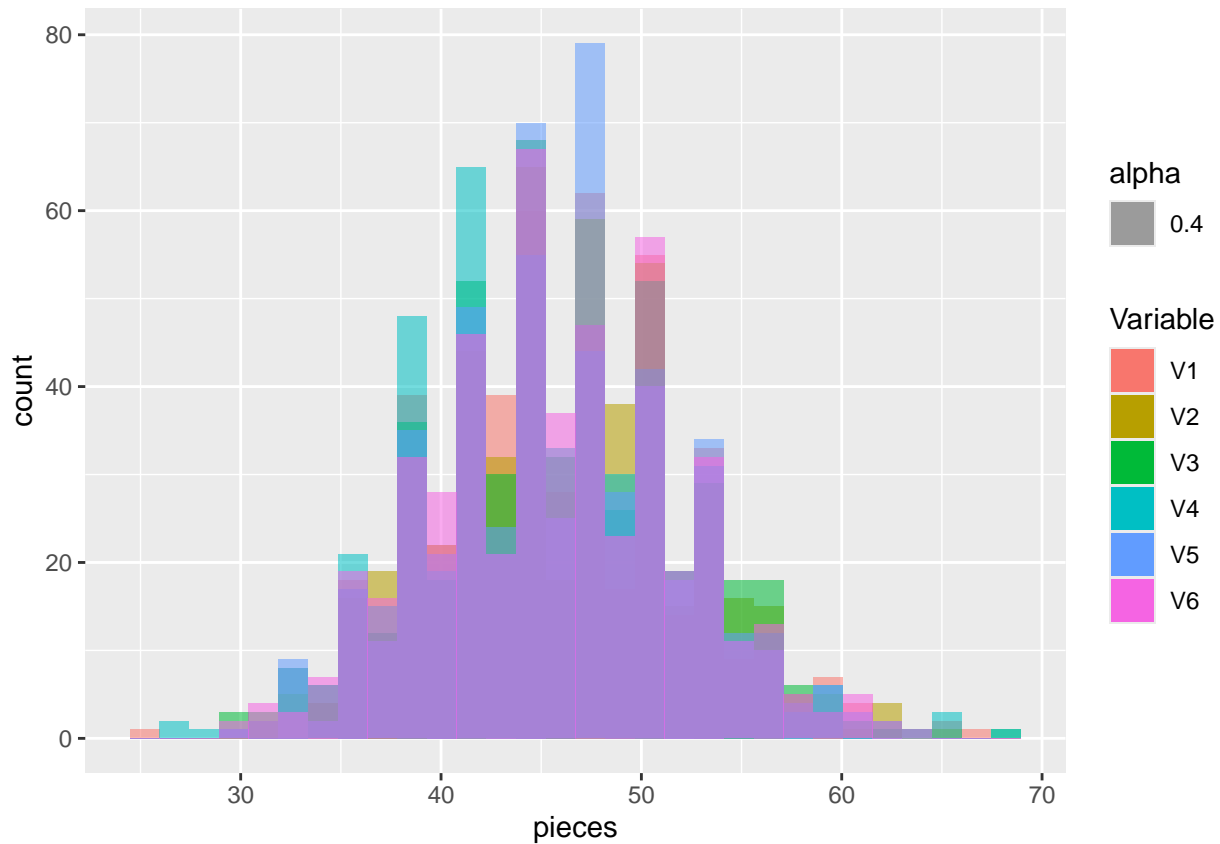
```

MnM_sample %>%
  pivot_longer(cols = 1:6, names_to = "Variable", values_to = "pieces") %>%

```

```
ggplot(aes(fill=Variable ,x= pieces, alpha= 0.4))+
  geom_histogram(position = "identity")
```

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



we can see that the variance distribution is Gamma like with shape and rate as seen below

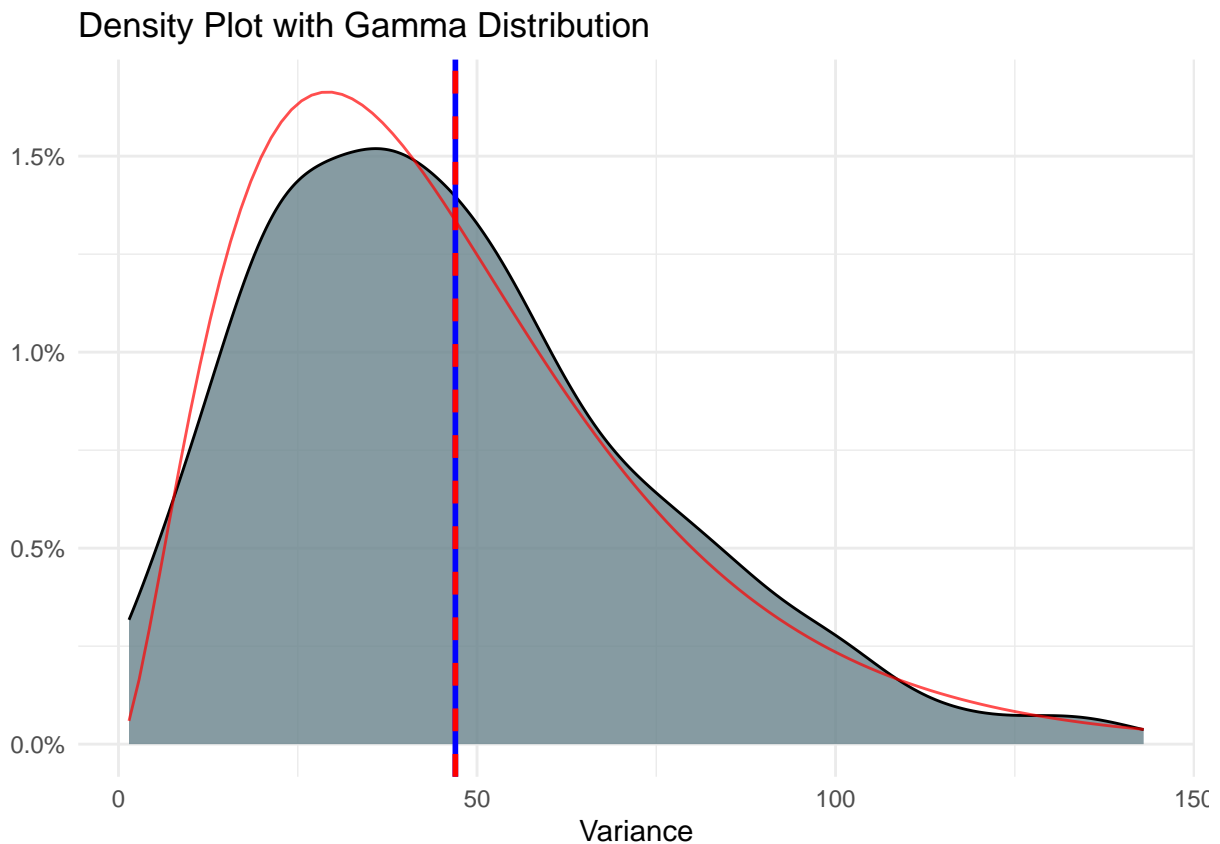
```
gamma_params <- fitdistr(MnM_sample$Variance, "gamma")$estimate
paste0("the parameters of the gamma shaped var is shape ", round(gamma_params[1],3)," and rate ", round
```

[1] "the parameters of the gamma shaped var is shape 2.635 and rate 0.056"

```
shape_est <- gamma_params["shape"]
scale_est <- 1 / gamma_params["rate"] #using Scale= 1/Rate

MnM_sample %>% #plot density
  ggplot(aes(x = Variance, fill = "orange")) +
  geom_density(fill= "lightblue4", alpha= 0.8) +
  geom_vline(xintercept = mean(MnM_sample$Variance), color = "blue", size=1) +
  geom_vline(xintercept = scale_est*shape_est, color = "red", linetype = "dashed", size=1) +
  stat_function(fun = dgamma, args = list(shape = shape_est, scale = scale_est),
    color = "red", alpha= 0.7) +
  labs(title = "Density Plot with Gamma Distribution",
    x = "Variance",
```

```
y = """) +
scale_y_continuous(label=scales::label_percent(.1)) +
theme_minimal()
```



#check too low color (under 10%) and sample by n number
 #use statistics to sample better low chance cases

optimizing best package

n*m types of snacks

We will create a function that create sample for each number of colors and package size we want, and then calculate some interesting parameters

```
mega_snack<- function(nn,n_unit,n_color)
{
  m_sample<- length(n_unit)*length(n_color)
  nul_mat= matrix(nrow = m_sample, ncol = 5)
  res<- cbind(rep(n_unit,length(n_color)),sort(rep(n_color,length(n_unit))),
              nul_mat)
  for (i in 1:(dim(res)[1]))
  {
    #print(c(res[i,1],res[i,2]))
  }
}
```



```

low_color<- 0.666*res[i,1]/(res[i,2])
small_sample<- sample_MnM(nn,res[i,1],res[i,2])
small_sample<-
  small_sample %>% as_data_frame() %>%
  mutate(even_count= rowSums(across(everything() , ~ .x %% 2 == 0))/n_color, #how many evens color
         even_evens= (rowSums(across(-c(even_count) , ~ .x %% 2 == 1)) %% 2 ==0)/n_color, #are the
         var_col=     apply(across(c(-even_count, even_evens)), 1, var), #var of candy per color/ ty
         all_even=     rowSums(across(everything() , ~ .x %% 2 == 0))== n_color,
         low_col=      rowSums(across(-c(even_count, even_evens,var_col,all_even), ~ .x <= low_color )
res[i,3]<- mean(small_sample$even_count)
res[i,4]<- mean(small_sample$even_evens)
res[i,5]<- mean(small_sample$var_col)
res[i,6]<- mean(small_sample$all_even)
res[i,7]<- mean(small_sample$low_col)
  }
colnames(res)<- c("n_unit", "n_color", "even_count", "even_evens", "var_col","all_even","low_color")
res
}

```

```

color_op<- 2:7
grams_op<- c(25,45,150,250,330,500,750,1000)
n_unit_op<- grams_op/gram

mega_snack_1<-
  mega_snack(500,n_unit_op,color_op) %>% as.data.frame() %>%
  mutate(n_unit= round(n_unit,1))

mega_snack_1 %>% head(6)

```

```

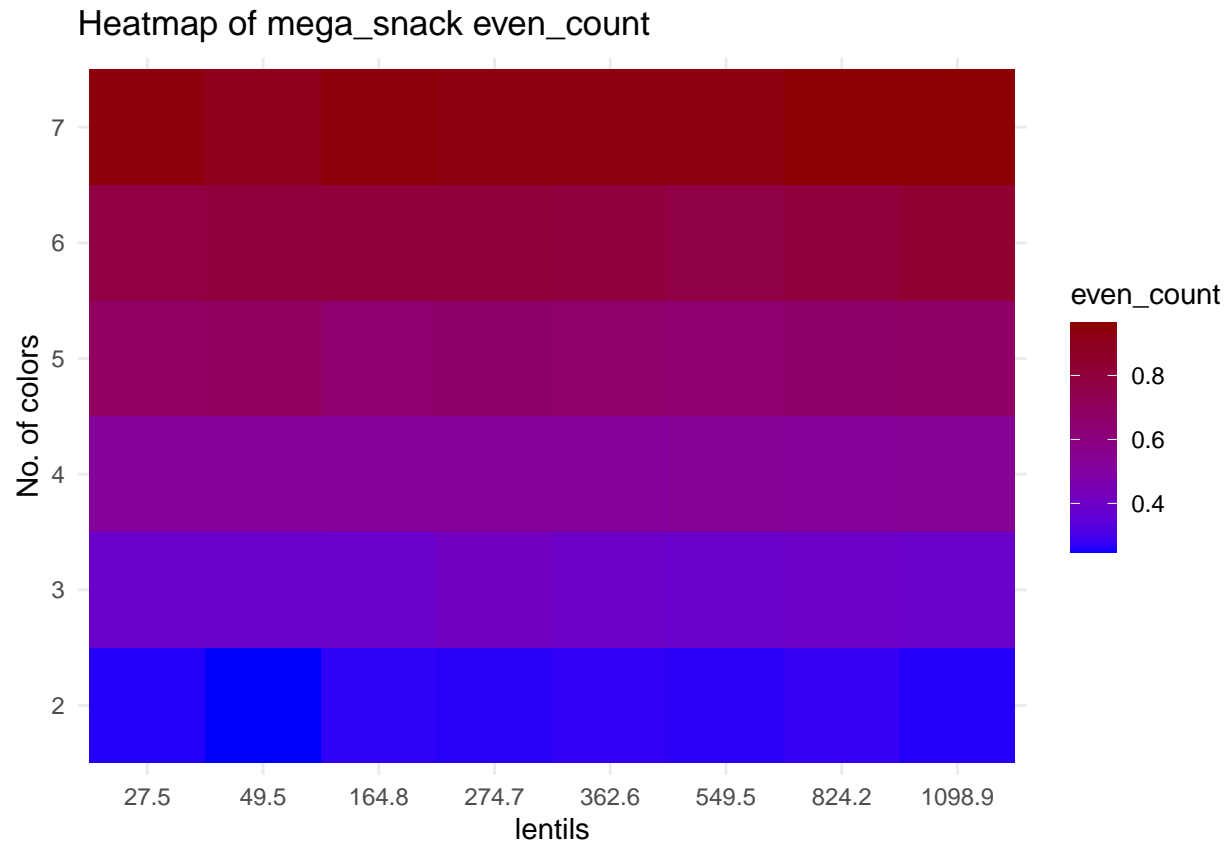
##   n_unit n_color even_count even_evens      var_col all_even low_color
## 1   27.5       2  0.2603810  0.1337190    69.54723   0.130   0.110
## 2   49.5       2  0.2475619  0.1440619   214.65000   0.108   0.026
## 3  164.8       2  0.2703476  0.1323905  2286.95159   0.128   0.000
## 4  274.7       2  0.2648952  0.1332143  6336.46057   0.130   0.000
## 5  362.6       2  0.2721095  0.1361714 11012.44906   0.122   0.000
## 6  549.5       2  0.2678000  0.1470333 25275.14270   0.124   0.000

```

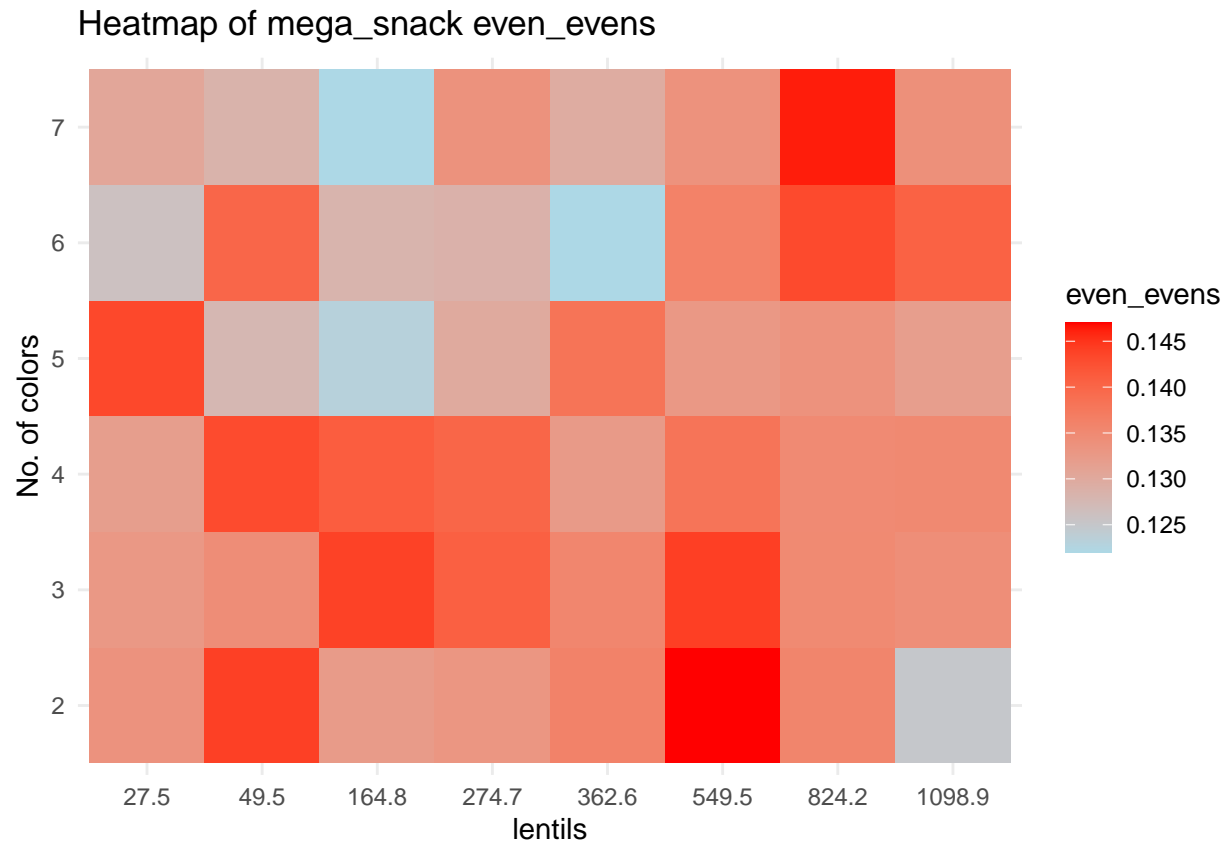
```

mega_snack_1 %>%
  ggplot(aes(x = factor((round( n_unit,1) )), y = factor(n_color ), fill = even_count )) +
  geom_tile() +
  scale_fill_gradient(low = "blue", high = "red4")+
  labs(title = "Heatmap of mega_snack even_count",
       x = "lentils",
       y = "No. of colors",
       fill = "even_count") +
  theme_minimal()

```

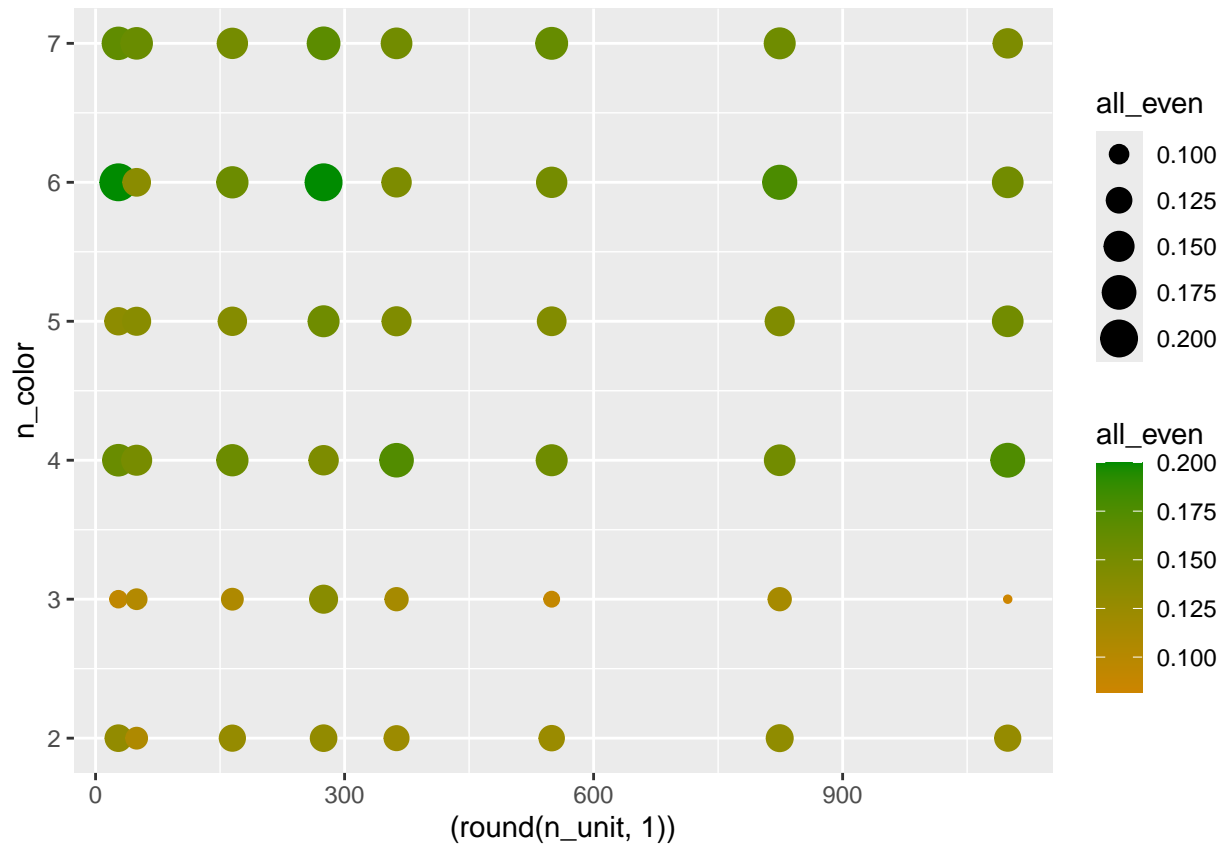


```
mega_snack_1 %>%
  ggplot(aes(x = factor((round( n_unit,1) )), y = factor(n_color ), fill = even_evens )) +
  geom_tile() +
  scale_fill_gradient(low = "lightblue", high = "red")+
  labs(title = "Heatmap of mega_snack even_evens",
       x = "lentils",
       y = "No. of colors",
       fill = "even_evens") +
  theme_minimal()
```



now let us see the probability of all even, and whether there is pattern.

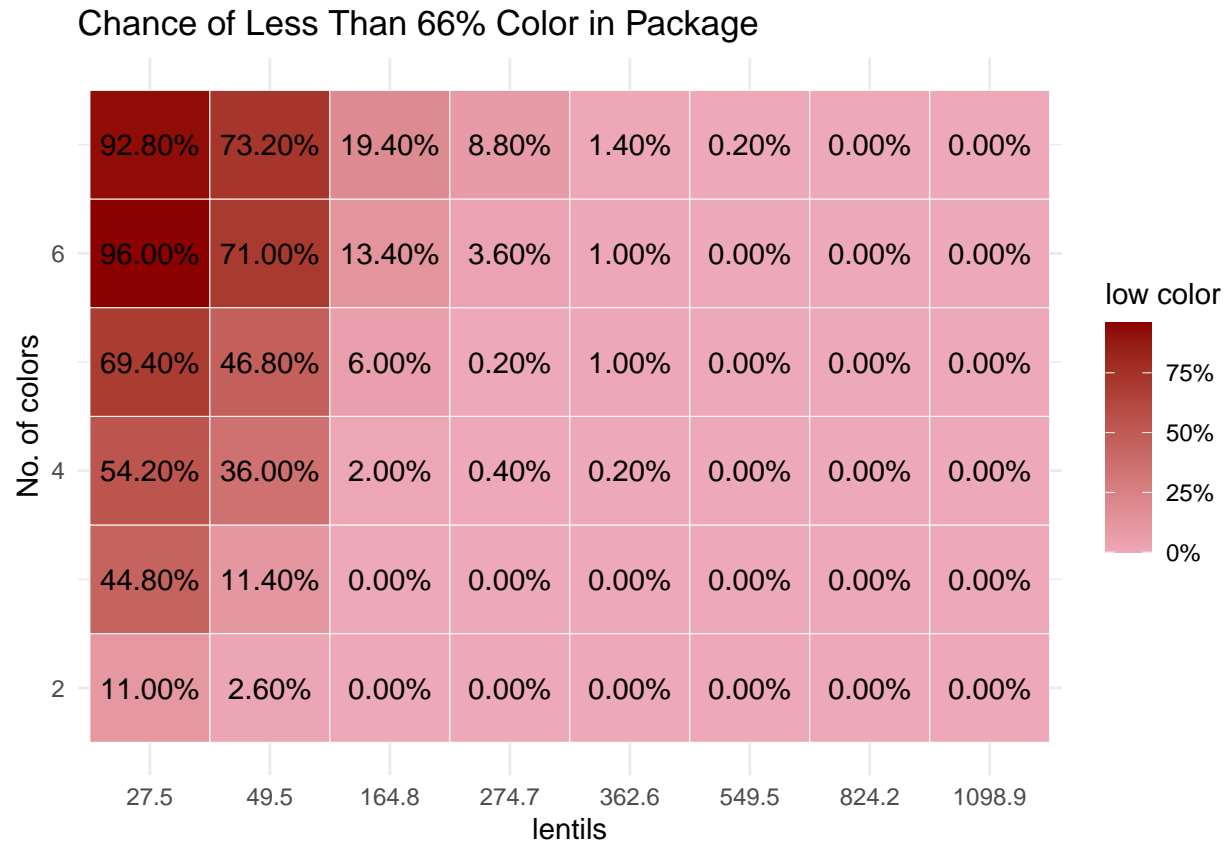
```
mega_snack_1 %>%
  ggplot(aes(x = (round( n_unit,1) ), y = n_color , color = all_even,size = all_even )) +
  geom_point() +
  scale_color_gradient(low = "orange3", high = "green4")
```



```
labs(title = "Heatmap of mega_snack all evens",
      x = "lentils",
      y = "No. of colors",
      color = "all_even") +
theme_minimal()
```

NULL

```
mega_snack_1 %>%
  ggplot(aes(x = factor(round(n_unit, 1)), y = n_color, fill = low_color)) +
  geom_tile(color = "white") +
  geom_text(aes(label = sprintf("%.2f%%", low_color * 100)), color = "black", size = 4) +
  scale_fill_gradient(low = "pink2", high = "red4", labels = scales::percent) +
  labs(title = "Chance of Less Than 66% Color in Package",
        x = "lentils",
        y = "No. of colors",
        fill = "low color") +
  theme_minimal()
```



As we can see, only the small package (less than 50 lentils) have high probability of at least one color to appear severely lower.

Therefore, splitting package by color on the big ones should be relatively even.