## 9\_real data example 2

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# Example with 50 counts (Real data from the U.S. Census Bureau Website)

## Generate 50 counts and compute the total count

The data table is retrieved from: https://www.census.gov/data/tables/2010/dec/2010-apportionment-data. html

Apportionment Population And Number of Representatives by State: 2010 Cencus.

```
d <- read.csv("ApportionmentPopulation2020_newdata.csv", header = F)</pre>
N \leftarrow sort(d$V4[2:51])
t <- sum(N)
N
                                         2
                                             2
                                                 2
                                                      2
                                                          2
                                                              3
                                                                                4
   [1]
          1
                            1
                                    1
                                                                   3
                                                                       3
                   1
                       1
                                1
                                                 7
## [18]
          4
               4
                   4
                       4
                            5
                                5
                                    5
                                         6
                                             6
                                                     7
                                                          8
                                                              8
                                                                   8
                                                                       8
                                                                           9
                                                                                9
## [35]
               9 10
                          12 13 14 14
                                           16 18
                      11
                                                   18 27
                                                             27
                                                                 36 53 435
## [1] 863
length(N)
## [1] 50
```

## Define alpha and sampling functions

```
# Define alpha based on the geometric mechanism
alpha <- 1/exp(1)

# pdf of the doubbe geometric distribution
probs <- function(n, k = 0){
    p <- c()
    for(i in 1:length(n)){
        p[i] <- alpha^(abs(n[i] - k))*(1-alpha)/(1 + alpha)
    }
    return(p)
}

# Chop the noise so that i is less than or equal to 50 (?)

# Set the first and the last probabilities(Boundaries) to adjust when using sample function
first_p = last_p = function(p){
    return(0.5*(1 - sum(p)))
}</pre>
```

```
# Define the function to sample noisy count (could be positive) from the dg distribution
samplenoise <- function(n, center = 0, i = 50){</pre>
  i_ = i-1
  return(sample(x = (-i + center): (i + center), size = n,
                prob = c(first_p(probs((-i_ + center):(i_ + center), center)),
                          probs((-i_ + center):(i_ + center), center),
                          last_p(probs((-i_ + center):(i_ + center), center)))))
}
# Define the function to sample posterior count (must be positive) from the dq distribution
samplepos <- function(n, center = 0, i = 50){</pre>
  # Two cases
  if (-i + center >= 0){
    result = samplenoise(n, center, i)
  } else {
    prob = probs(0:(i + center), center)
    result = sample(x = 0:(i + center), size = n, prob = prob / sum(prob))
 return(result)
}
# Define the function to find the pi in multinomial idea
pi <- function(pos){</pre>
  p \leftarrow c()
  for(i in 1:length(pos)){
    p[i] <- pos[i] / sum(pos)</pre>
  }
  return(p)
}
```

## Find noisy counts and the posterior of the total and individuals

All algorithms share the same noisy counts.

- 1. Algorithm 1: New total and new components for each try
- 2. Algorithm 2: New total with fixed components
- 3. Algorithm 3: Only use noise counts (Not Bayesian). If noisy count is negative, use the posterior mode.

```
# Generate n counts
n = 10000

# Create vectors for results
p1_total <- c()
p2_total <- c()
t3_noise <- c()
p1_N <- matrix(data = NA, n, 50)
N3_noise <- matrix(data = NA, n, 50)
for(i in 1:n){</pre>
```

```
# Noisy count of the total
  t.noise <- samplenoise(1, center = t, i = 50)</pre>
  # Algorithm 1
    ## Posterior count of the total
  p1_total[i] <- samplepos(1, center = t.noise, i = 50)</pre>
    ## Sample the individual noisy counts
  N_noise <- c()
  for(j in 1:length(N)){
    N_{\text{noise}}[j] \leftarrow \text{samplenoise}(1, \text{center} = N[j], i = 50)
    ## Sample the individual posterior counts
  posterior_N <- c()</pre>
  for(k in 1:length(N_noise)){
    posterior_N[k] <- samplepos(n = 1, center = N_noise[k], i = 50)</pre>
  p1_N[i,] <- posterior_N</pre>
  # Algorithm 2
  p2_total[i] <- p1_total[i]</pre>
  # Algorithm 3
  t3_noise[i] <- t.noise
  N3_noise[i,] <- N_noise
# Algorithm 3 Adjustments
# Function used to find the posterior mode
Mode <- function(x) {</pre>
  ux <- unique(x)
  ux[which.max(tabulate(match(x, ux)))]
# Substitute negative noisy counts with the posterior mode of that state
for(i in 1:n){
  for(j in 1:50){
    if(N3_noise[i,j] < 0){</pre>
      N3\_noise[i,j] \leftarrow Mode(p1\_N[,j])
    }
  }
}
```

## Multinomial idea

Generate each individual count of multinomial idea

```
set.seed(1)
multi_1 <- matrix(data = NA, 50, n)</pre>
```

```
multi_2 <- matrix(data = NA, 50, n)
multi_3 <- matrix(data = NA, 50, n)

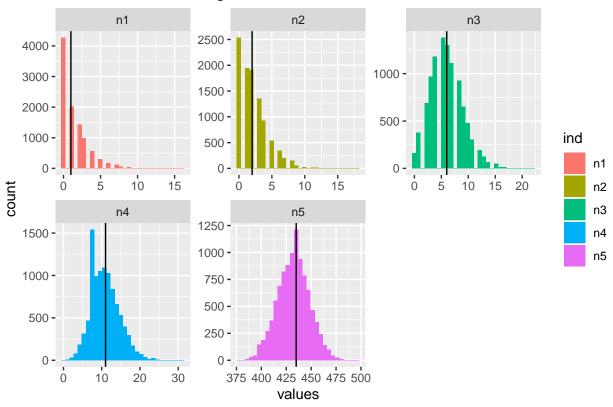
for(i in 1:n){
    # Algorithm 1
    multi_1[, i] <- rmultinom(n = 1, size = p1_total[i], prob = pi(p1_N[i,]))

# Algorithm 2: Probabilities are the same for all 10000 runs.
multi_2[, i] <- rmultinom(n = 1, size = p2_total[i], prob = pi(p1_N[1,]))

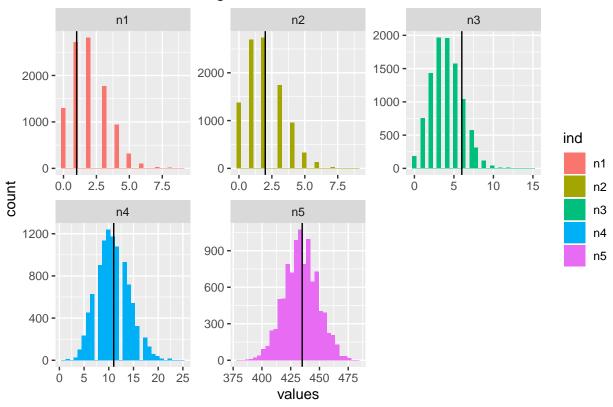
# Algorithm 3: Only use noisy counts for each run
multi_3[, i] <- rmultinom(n = 1, size = t3_noise[i], prob = pi(N3_noise[i,]))
}</pre>
```

#### Plot the results

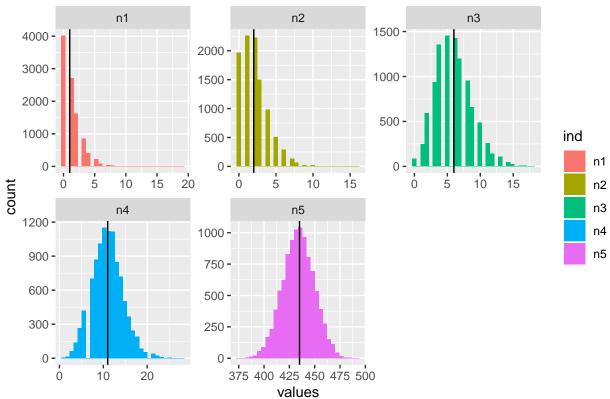
## Multinomial idea - Algorithm 1



## Multinomial idea - Algorithm 2



## Multinomial idea – Algorithm 3



## Comparison

## Variance

Table 1: Variances with the multinomial idea

	n1	n2	n3	n4	n5
Algorithm 1 Algorithm 2 Algorithm 3	1.9179	1.9771	3.9436	10.5247	214.3874

#### Bias

```
 \begin{aligned} df\_bias &\leftarrow data.frame(n1 = c(bias(result1\$n1, rep(N[1], n)), bias(result2\$n1, rep(N[1], n)), \\ & bias(result3\$n1, rep(N[1], n))), \\ & n2 = c(bias(result1\$n2, rep(N[12], n)), bias(result2\$n2, rep(N[12], n)), \end{aligned}
```

Table 2: Bias with the multinomial idea

	n1	n2	n3	n4	n5
Algorithm 1	0.4782	0.2022	-0.0435	-0.0526	-2.6779
Algorithm 2	0.9918	-0.0100	-2.0165	-0.1130	-0.9524
Algorithm 3	0.2476	0.1554	-0.0413	-0.0162	-1.3737

#### Mean Squared Error

Table 3: MSE with the multinomial idea

	n1	n2	n3	n4	n5
Algorithm 1 Algorithm 2 Algorithm 3	2.9014	1.9770	8.0095	10.5364	215.2730