

```

[1] "# of discriminatory genes = 5"

[1] "fold change for discriminatory genes: 1"

[1] "fold change for nondiscriminatory genes: 0.1"

choose_k
  2
100

```

The optimal number of clusters K is set as the most frequently found K from the 100 simulations tabulated above. It is found by using the BIC criterion after running the unpenalized EM algorithm on K spanning from 2 to 8.

Using the last set of simulated counts, I ran a grid search across varying tuning parameters. As done in Pan et al, I fixed $\lambda_1 = 1$, and searched over $\lambda_2 = (0.1, 0.2, \dots, 2)$ and $\tau = (0.1, 0.2, \dots, 2)$:

```

[1] "lambda1, lambda2, tau, BIC:"

      [,1] [,2] [,3]      [,4]
[1,]    1  0.3  0.2 19149.87
[2,]    1  0.3  0.3 19149.87
[3,]    1  0.3  0.4 19149.87
[4,]    1  0.3  0.5 19149.87
[5,]    1  0.3  0.6 19149.87
[6,]    1  0.3  0.7 19149.87
[7,]    1  0.3  0.8 19149.87
[8,]    1  0.3  0.9 19149.87
[9,]    1  0.3  1.0 19149.87
[10,]   1  0.3  1.1 19149.87
[11,]   1  0.3  1.2 19149.87
[12,]   1  0.3  1.3 19149.87
[13,]   1  0.3  1.4 19149.87
[14,]   1  0.4  0.1 19149.87
[15,]   1  0.4  0.2 19149.87
[16,]   1  0.4  0.3 19149.87
[17,]   1  0.4  0.4 19149.87
[18,]   1  0.4  0.5 19149.87
[19,]   1  0.4  0.6 19149.87
[20,]   1  0.4  0.7 19149.87
[21,]   1  0.4  0.8 19149.87
[22,]   1  0.4  0.9 19149.87
[23,]   1  0.4  1.0 19149.87
[24,]   1  0.4  1.1 19149.87
[25,]   1  0.4  1.2 19149.87
[26,]   1  0.4  1.3 19149.87
[27,]   1  0.4  1.4 19149.87

```

[28,]	1	0.5	0.1	19149.87
[29,]	1	0.5	0.2	19149.87
[30,]	1	0.5	0.3	19149.87
[31,]	1	0.5	0.4	19149.87
[32,]	1	0.5	0.5	19149.87
[33,]	1	0.5	0.6	19149.87
[34,]	1	0.5	0.7	19149.87
[35,]	1	0.5	0.8	19149.87
[36,]	1	0.5	0.9	19149.87
[37,]	1	0.5	1.0	19149.87
[38,]	1	0.5	1.1	19149.87
[39,]	1	0.5	1.2	19149.87
[40,]	1	0.5	1.3	19149.87
[41,]	1	0.6	0.1	19149.87
[42,]	1	0.6	0.2	19149.87
[43,]	1	0.6	0.3	19149.87
[44,]	1	0.6	0.4	19149.87
[45,]	1	0.6	0.5	19149.87
[46,]	1	0.6	0.6	19149.87
[47,]	1	0.6	0.7	19149.87
[48,]	1	0.6	0.8	19149.87
[49,]	1	0.6	0.9	19149.87
[50,]	1	0.6	1.0	19149.87
[51,]	1	0.6	1.1	19149.87
[52,]	1	0.6	1.2	19149.87
[53,]	1	0.7	0.1	19149.87
[54,]	1	0.7	0.2	19149.87
[55,]	1	0.7	0.3	19149.87
[56,]	1	0.7	0.4	19149.87
[57,]	1	0.7	0.5	19149.87
[58,]	1	0.7	0.6	19149.87
[59,]	1	0.7	0.7	19149.87
[60,]	1	0.7	0.8	19149.87
[61,]	1	0.7	0.9	19149.87
[62,]	1	0.7	1.0	19149.87
[63,]	1	0.7	1.1	19149.87
[64,]	1	0.8	0.1	19149.87
[65,]	1	0.8	0.2	19149.87
[66,]	1	0.8	0.3	19149.87
[67,]	1	0.8	0.4	19149.87
[68,]	1	0.8	0.5	19149.87
[69,]	1	0.8	0.6	19149.87
[70,]	1	0.8	0.7	19149.87
[71,]	1	0.8	0.8	19149.87
[72,]	1	0.8	0.9	19149.87
[73,]	1	0.8	1.0	19149.87

[74,]	1	0.9	0.1	19149.87
[75,]	1	0.9	0.2	19149.87
[76,]	1	0.9	0.3	19149.87
[77,]	1	0.9	0.4	19149.87
[78,]	1	0.9	0.5	19149.87
[79,]	1	0.9	0.6	19149.87
[80,]	1	0.9	0.7	19149.87
[81,]	1	0.9	0.8	19149.87
[82,]	1	0.9	0.9	19149.87
[83,]	1	1.0	0.1	19149.87
[84,]	1	1.0	0.2	19149.87
[85,]	1	1.0	0.3	19149.87
[86,]	1	1.0	0.4	19149.87
[87,]	1	1.0	0.5	19149.87
[88,]	1	1.0	0.6	19149.87
[89,]	1	1.0	0.7	19149.87
[90,]	1	1.0	0.8	19149.87
[91,]	1	1.1	0.1	19149.87
[92,]	1	1.1	0.2	19149.87
[93,]	1	1.1	0.3	19149.87
[94,]	1	1.1	0.4	19149.87
[95,]	1	1.1	0.5	19149.87
[96,]	1	1.1	0.6	19149.87
[97,]	1	1.1	0.7	19149.87
[98,]	1	1.2	0.1	19149.87
[99,]	1	1.2	0.2	19149.87
[100,]	1	1.2	0.3	19149.87
[101,]	1	1.2	0.4	19149.87
[102,]	1	1.2	0.5	19149.87
[103,]	1	1.2	0.6	19149.87
[104,]	1	1.3	0.1	19149.87
[105,]	1	1.3	0.2	19149.87
[106,]	1	1.3	0.3	19149.87
[107,]	1	1.3	0.4	19149.87
[108,]	1	1.3	0.5	19149.87
[109,]	1	1.4	0.1	19149.87
[110,]	1	1.4	0.2	19149.87
[111,]	1	1.4	0.3	19149.87
[112,]	1	1.4	0.4	19149.87
[113,]	1	1.5	0.1	19149.87
[114,]	1	1.5	0.2	19149.87
[115,]	1	1.5	0.3	19149.87
[116,]	1	1.6	0.1	19149.87
[117,]	1	1.6	0.2	19149.87
[118,]	1	1.7	0.1	19149.87

The results of the final run based on optimal tuning parameters are below:
Below are the summary of results:

```
[1] "Mean pi: 0.608695652173913" "Mean pi: 0.391304347826087"
```

```
[1] "First 3 genes:"
```

```
      [,1]      [,2]
[1,] 2.888701 4.842502
[2,] 4.103085 6.089145
[3,] 3.616875 5.600119
```

```
[1] "Last 3 genes:"
```

```
      [,1]      [,2]
[98,] 6.312426 6.485494
[99,] 5.867213 6.030781
[100,] 6.292918 6.463273
```

```
[1] "Mean % of nondiscriminatory genes: 0.9499"
```

```
[1] "Mean ARI: 1"
```

```
[1] "Mean sensitivity: 1"
```

```
[1] "Mean false positive rate: 0.000105263157894737"
```