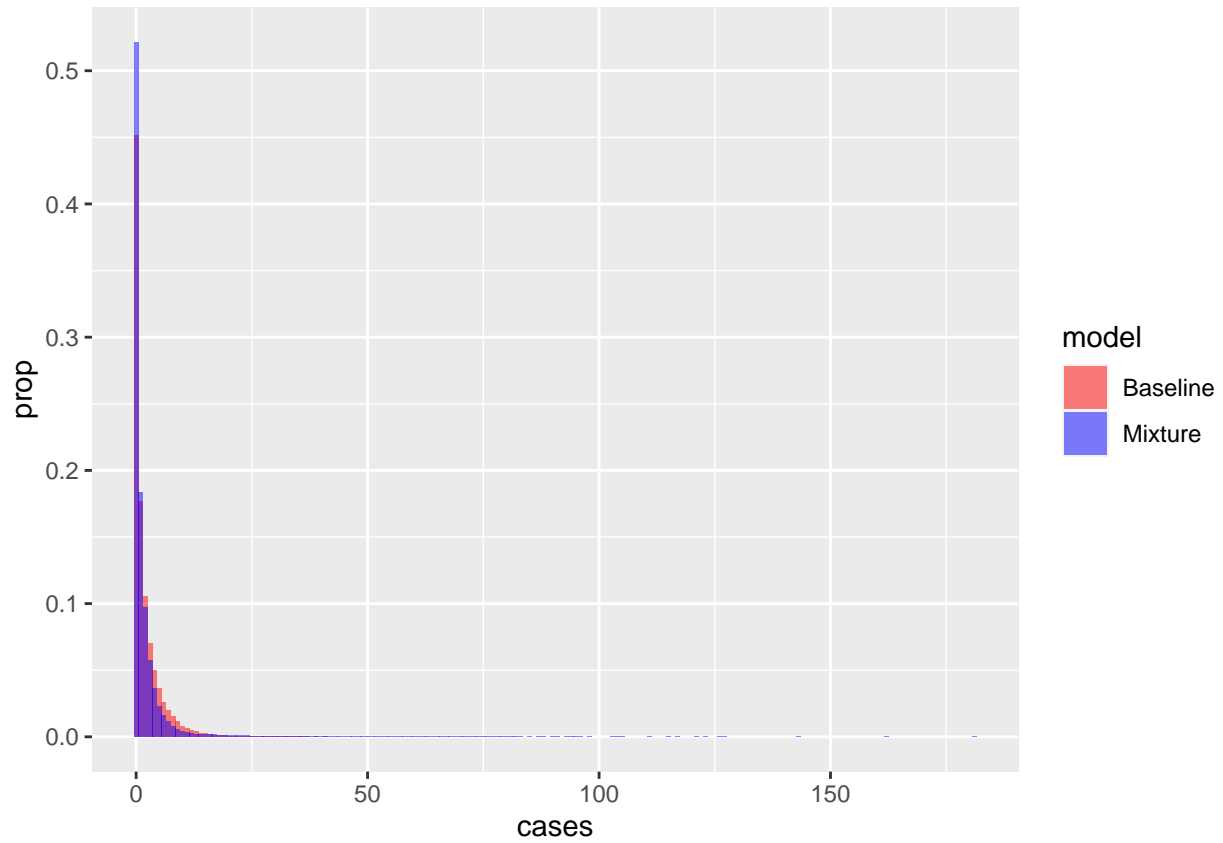


## Superspreading negative binomial simulations



##	cases	n
## 1	0	52155
## 2	1	18382
## 3	2	9737
## 4	3	5745
## 5	4	3644
## 6	5	2320
## 7	6	1596
## 8	7	1130
## 9	8	780
## 10	9	581
## 11	10	426
## 12	11	320
## 13	12	275
## 14	13	224
## 15	14	204
## 16	15	194

## 17	16	177
## 18	17	173
## 19	18	134
## 20	19	114
## 21	20	122
## 22	21	103
## 23	22	88
## 24	23	85
## 25	24	86
## 26	25	73
## 27	26	69
## 28	27	62
## 29	28	61
## 30	29	45
## 31	30	56
## 32	31	50
## 33	32	51
## 34	33	41
## 35	34	50
## 36	35	48
## 37	36	31
## 38	37	46
## 39	38	34
## 40	39	39
## 41	40	18
## 42	41	16
## 43	42	25
## 44	43	19
## 45	44	13
## 46	45	20
## 47	46	15
## 48	47	13
## 49	48	18
## 50	49	17
## 51	50	22
## 52	51	16
## 53	52	12
## 54	53	12
## 55	54	12
## 56	55	12
## 57	56	9
## 58	57	13
## 59	58	9
## 60	59	3
## 61	60	6
## 62	61	6
## 63	62	9
## 64	63	4
## 65	64	9
## 66	65	8
## 67	66	6
## 68	67	8
## 69	68	2
## 70	69	6

```
## 71      70      10
## 72      71       5
## 73      72      10
## 74      73       2
## 75      74       4
## 76      75       5
## 77      76       3
## 78      77       1
## 79      78       3
## 80      79       4
## 81      80       2
## 82      81       3
## 83      82       2
## 84      83       4
## 85      85       2
## 86      87       4
## 87      88       1
## 88      90       2
## 89      91       5
## 90      93       1
## 91      94       1
## 92      95       1
## 93      96       1
## 94      98       1
## 95     103       1
## 96     104       1
## 97     105       1
## 98     111       1
## 99     115       1
## 100     117       1
## 101     121       1
## 102     123       1
## 103     126       1
## 104     127       2
## 105     143       1
## 106     162       1
## 107     181       1
```

If  $p$  is small but  $\delta$  is large, then the range of cases is large.

Now we wish to plot typical chains.

First lets look at the mean size of small (clusters  $< 10$ ) and large clusters ( $> 10$ ). The median clusters are typically bigger in the standard model than the mixture model but the mean is smaller, and mean chain sizes conditioned on extinction are similar in both models. If chopping the chain should condition on extinction/non-extinction?

```
## [1] 8461.335      10.000 144973.000
```

```
## [1] 2504
```

```
## [1] 1.709763
```

```
## [1] 5526.246      10.000 38535.000
```

```
## [1] 3389

## [1] 1.666116

## [1] 57.73043

## [1] 30

## [1] 1.824859

## [1] 20.05

## [1] 16.5

## [1] 1.666667
```

```
##      [,1] [,2] [,3] [,4] [,5] [,6] [,7] [,8] [,9] [,10]
## [1,]    1    1    1    1    1    1    1    1    1    1
## [2,]    1    2    0    5    0    0    4    1    0    1
## [3,]    1    5    0    6    0    0    8    2    0    0
## [4,]    0    2    0   14    0    0    4    1    0    0
## [5,]    0    0    0   13    0    0    1    2    0    0
## [6,]    0    0    0   16    0    0    0    2    0    0
## [7,]    0    0    0   15    0    0    0    2    0    0
## [8,]    0    0    0   13    0    0    0    3    0    0
## [9,]    0    0    0   20    0    0    0    1    0    0
## [10,]   0    0    0    7    0    0    0    0    0    0
## [11,]   0    0    0   11    0    0    0    0    0    0
## [12,]   0    0    0    7    0    0    0    0    0    0
## [13,]   0    0    0    2    0    0    0    0    0    0
## [14,]   0    0    0    1    0    0    0    0    0    0
## [15,]   0    0    0    0    0    0    0    0    0    0
## [16,]   0    0    0    0    0    0    0    0    0    0
## [17,]   0    0    0    0    0    0    0    0    0    0
## [18,]   0    0    0    0    0    0    0    0    0    0
## [19,]   0    0    0    0    0    0    0    0    0    0
## [20,]   0    0    0    0    0    0    0    0    0    0
## [21,]   0    0    0    0    0    0    0    0    0    0
```

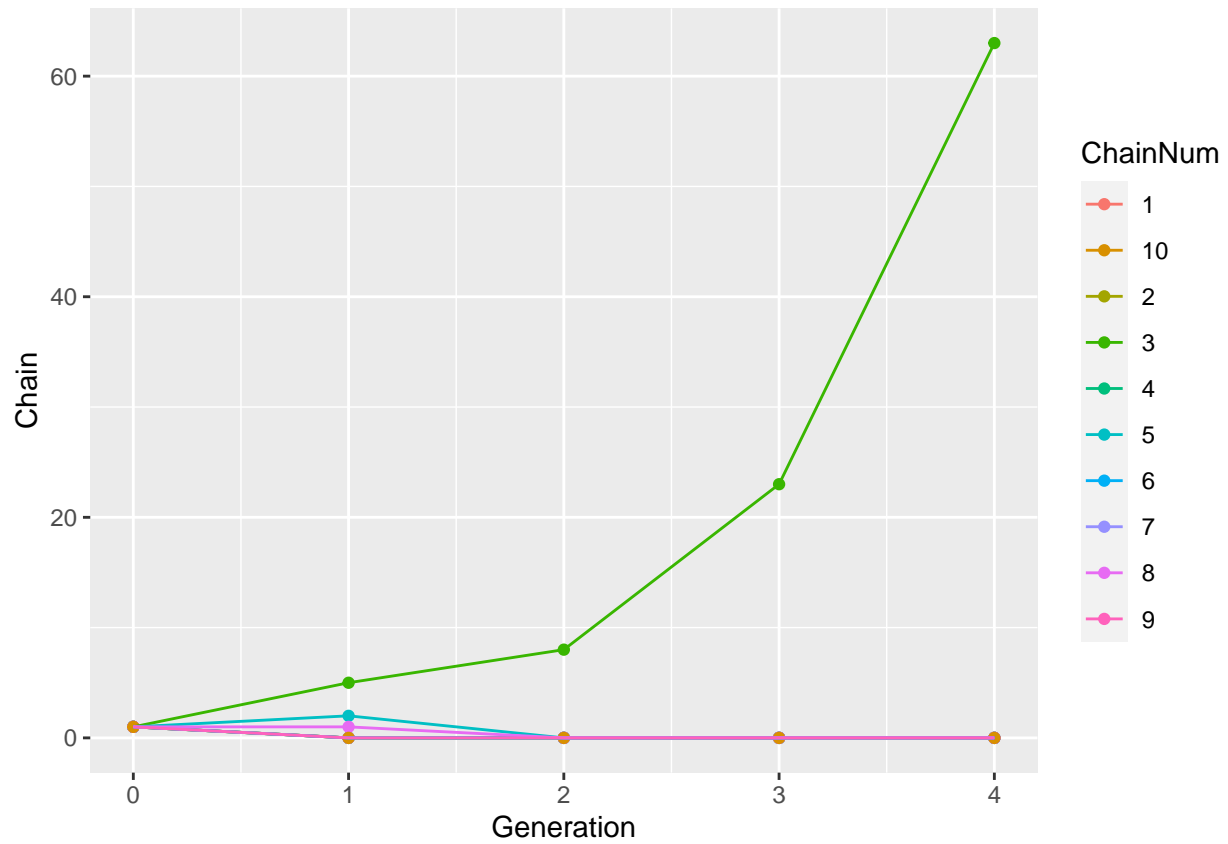
Probability of a major outbreak for mixture model is about 0.1.

```
##      [,1] [,2] [,3] [,4] [,5] [,6] [,7] [,8] [,9] [,10]
## [1,]    1    1    1    1    1    1    1    1    1    1
## [2,]    0    0    5    0    2    0    0    1    0    0
## [3,]    0    0    8    0    0    0    0    0    0    0
## [4,]    0    0   23    0    0    0    0    0    0    0
## [5,]    0    0   63    0    0    0    0    0    0    0
## [6,]    0    0  299    0    0    0    0    0    0    0
## [7,]    0    0  773    0    0    0    0    0    0    0
## [8,]    0    0 1247    0    0    0    0    0    0    0
## [9,]    0    0 2503    0    0    0    0    0    0    0
## [10,]   0    0 4698    0    0    0    0    0    0    0
## [11,]   0    0 9545    0    0    0    0    0    0    0
```

Probability of a major outbreak for standard model is about 0.4. Here four have taken off, but none are as large as the mixture that has taken off?

##		[,1]	[,2]	[,3]	[,4]	[,5]	[,6]	[,7]	[,8]	[,9]	[,10]
##	[1,]	1	1	1	1	1	1	1	1	1	1
##	[2,]	0	0	0	0	1	0	0	0	6	0
##	[3,]	0	0	0	0	3	0	0	0	13	0
##	[4,]	0	0	0	0	9	0	0	0	22	0
##	[5,]	0	0	0	0	5	0	0	0	64	0
##	[6,]	0	0	0	0	8	0	0	0	111	0
##	[7,]	0	0	0	0	17	0	0	0	243	0
##	[8,]	0	0	0	0	41	0	0	0	499	0
##	[9,]	0	0	0	0	61	0	0	0	1022	0
##	[10,]	0	0	0	0	120	0	0	0	2015	0
##	[11,]	0	0	0	0	279	0	0	0	3946	0

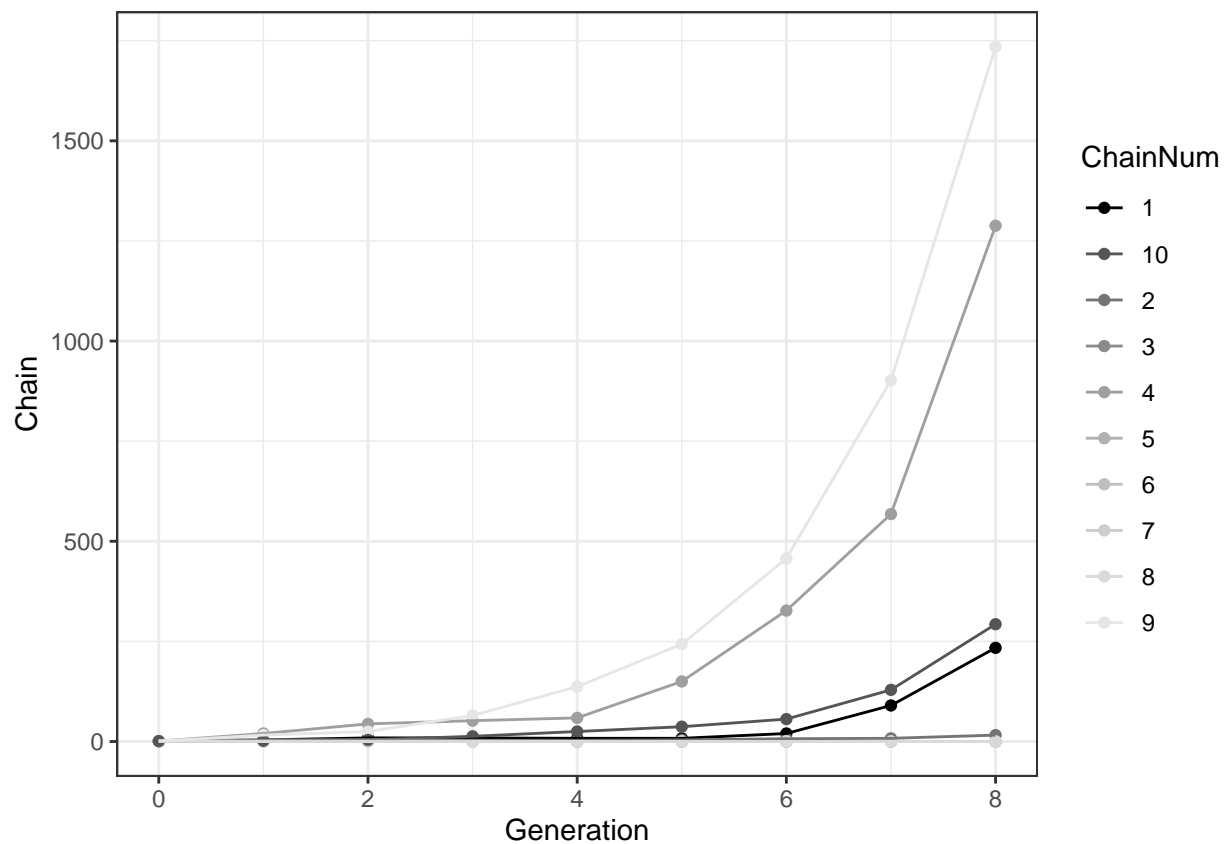
Plot the mixture chains assuming  $k=1/2$ :

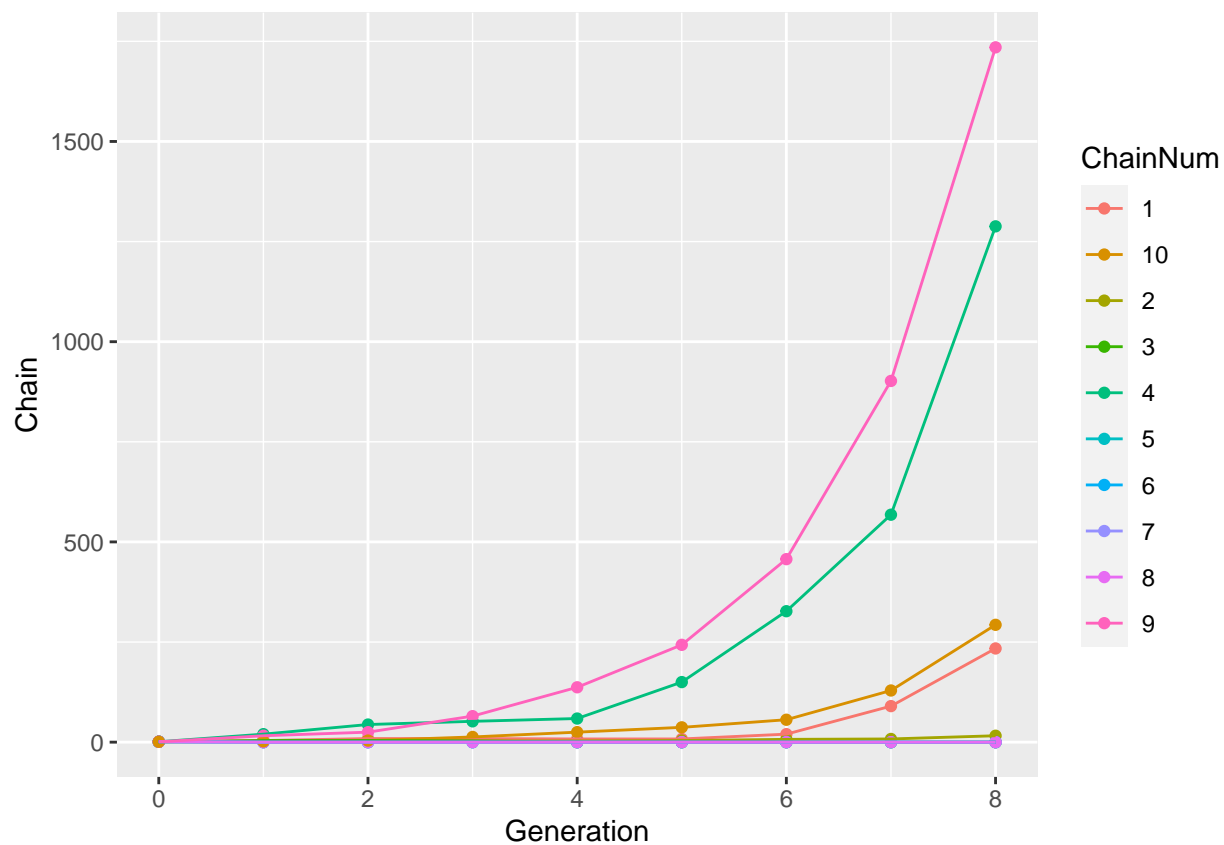


##		[,1]	[,2]	[,3]	[,4]	[,5]	[,6]	[,7]	[,8]	[,9]	[,10]
##	[1,]	1	1	1	1	1	1	1	1	1	1
##	[2,]	4	1	4	20	0	0	0	0	16	2
##	[3,]	9	2	4	44	0	0	0	0	25	4
##	[4,]	9	3	3	52	0	0	0	0	65	13
##	[5,]	8	3	0	59	0	0	0	0	137	25
##	[6,]	8	4	0	150	0	0	0	0	243	37
##	[7,]	20	7	0	327	0	0	0	0	457	56

```
## [8,] 90 8 0 568 0 0 0 0 902 129
## [9,] 234 16 0 1288 0 0 0 0 1735 293
## [10,] 412 22 0 2446 0 0 0 0 4001 589
## [11,] 948 45 0 4897 0 0 0 0 8064 1228
```

Plot the mixture chains assuming  $k=2$ :





##	[,1]	[,2]	[,3]	[,4]	[,5]	[,6]	[,7]	[,8]	[,9]	[,10]
## [1,]	1	1	1	1	1	1	1	1	1	1
## [2,]	5	3	7	0	4	0	0	1	5	0
## [3,]	9	8	17	0	7	0	0	1	21	0
## [4,]	17	27	28	0	8	0	0	3	47	0
## [5,]	22	64	46	0	20	0	0	2	107	0
## [6,]	39	123	103	0	35	0	0	6	197	0
## [7,]	91	270	195	0	64	0	0	11	407	0
## [8,]	172	541	352	0	107	0	0	18	831	0
## [9,]	322	1085	686	0	165	0	0	34	1571	0
## [10,]	667	2250	1309	0	278	0	0	77	3198	0
## [11,]	1299	4474	2482	0	542	0	0	174	6234	0

Median generation time same for all mixtures. Generation time has most variability for  $p=0.05$ , with 25% taking 11 generations or more to take off. 25% take 3 generations or less (most explosive for  $p=0.05, 0.1$ ). 75%

take 9 generations or less if  $p=0.1$  (most explosive?)

