X1 X2 p model **ZD Mutation Sampler** 1 1 1 0.09765625 causal strengths 1 1 0 0.05859375 **X1 X2** 0 0.0234375 1 1 0.5 0 0 0 0 0.0703125 1 0 0 0.5 0 1 1 0.1171875 0 0 0 1 0.0703125 0 0 bias 0.5 0.140625 0 0 1 baserates 0.25 0.25 0.25 0 0 0 0.421875

| | (| |
|--|---|--|
| | | |

10000

Input parameters (black lines)

nChains

betavar

meanChainlen

13

- **X1** 1 1

1

1 0

0

0

0

Υ 1

1

0

0

1

1

0

0

mean sim joint distr

X2

1

0

1

0

1

0

1

0

input joint distr



0.147876923075515 0.0592923076963512

0.0616769230808199

0.11971538461571

0.0641076923114395

0.122576923077072

0.400815384598411

р

- 0.0239384615446807

X1 X2 p model IK BayesianMS 0.09765625 causal strengths 0.05859375 **X1 X2** 0.0234375 0.5 0.0703125 0.5 0.1171875 0.0703125 bias 0.5 0.140625 baserates 0.25 0.25 0.25 0.421875 mean sim joint distr state **X1** Υ **X2** р nChains 0.14777692 0.05993846

| betavar | |
|---------|--|
| | |

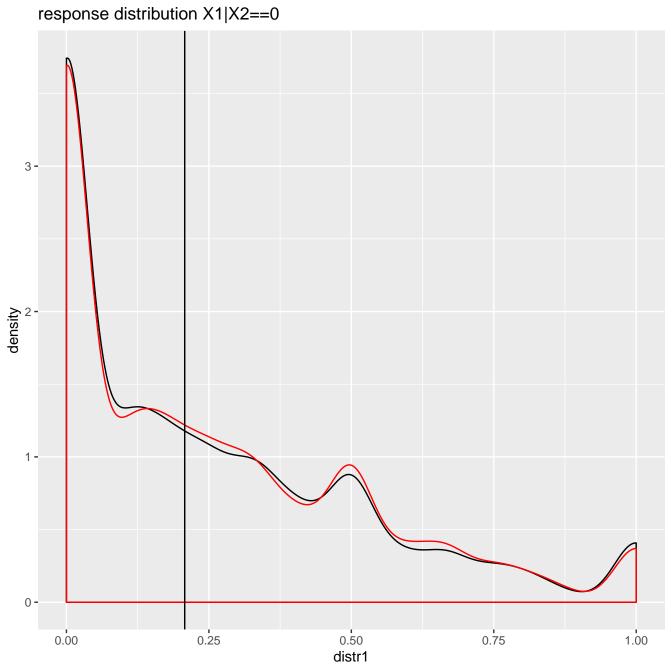
meanChainlen

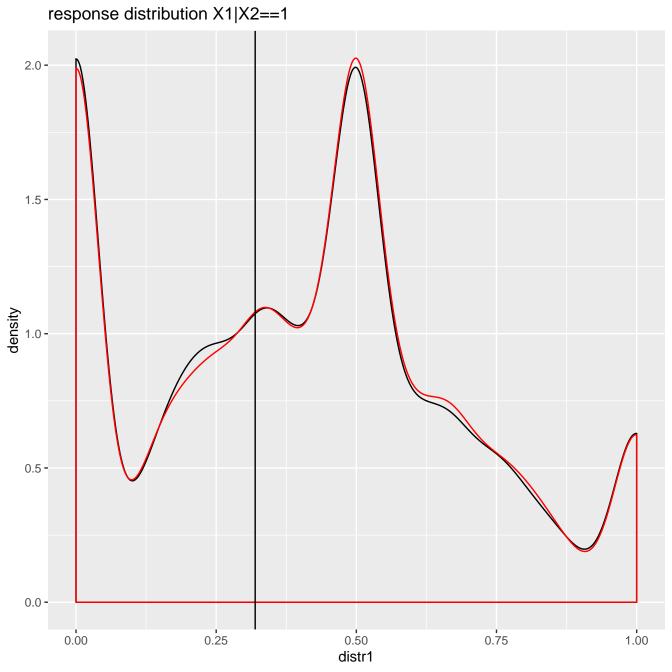
Input parameters (red lines)

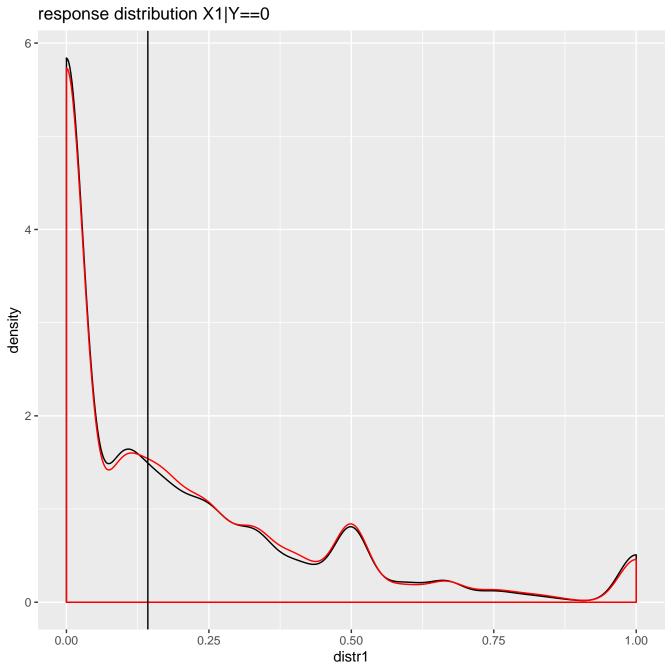
input joint distr

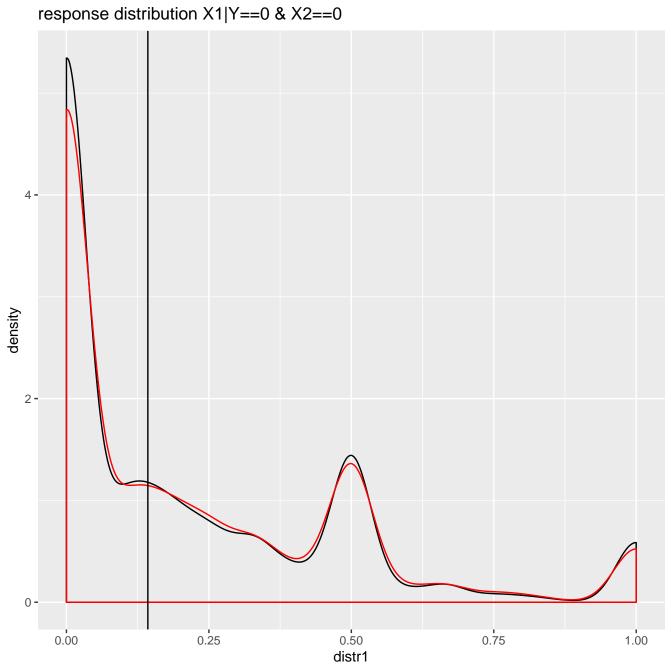
- 0.02432308 0.06381538 0.11762308 0.06343077 0.12037692

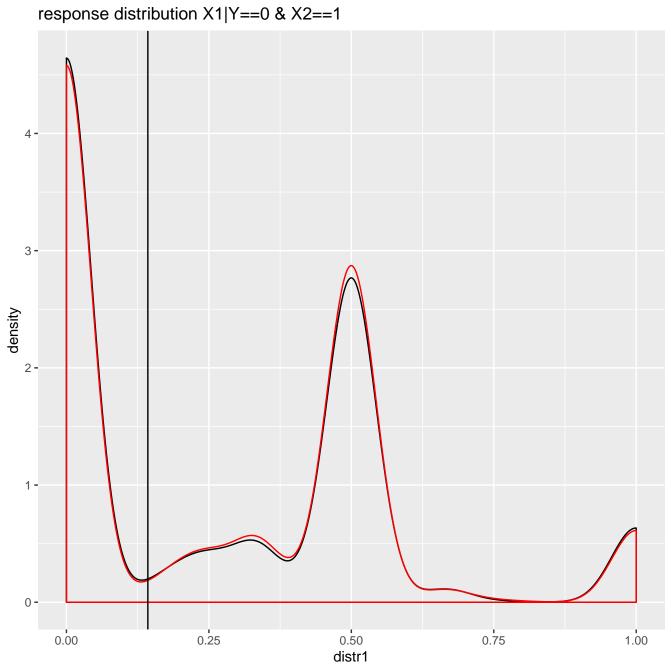
0.40271538

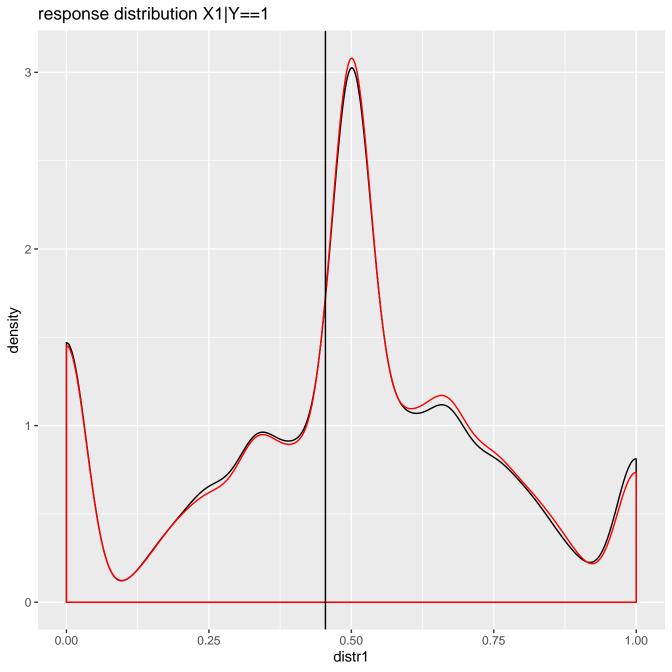




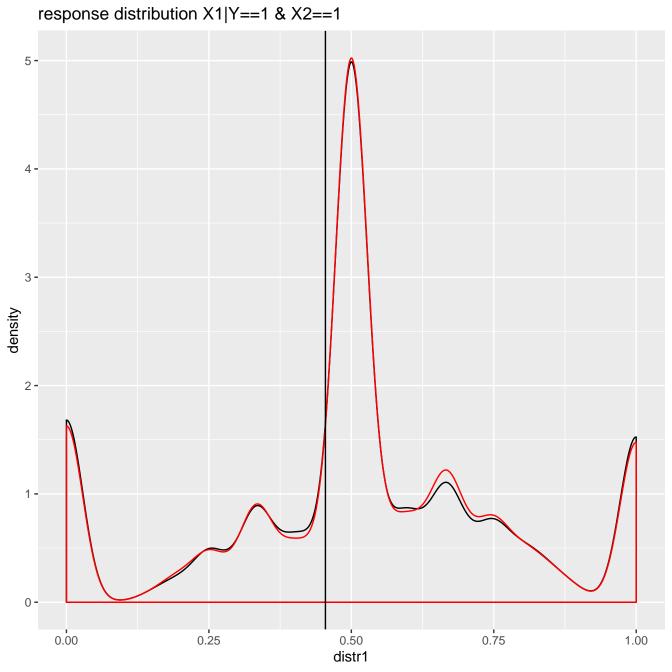


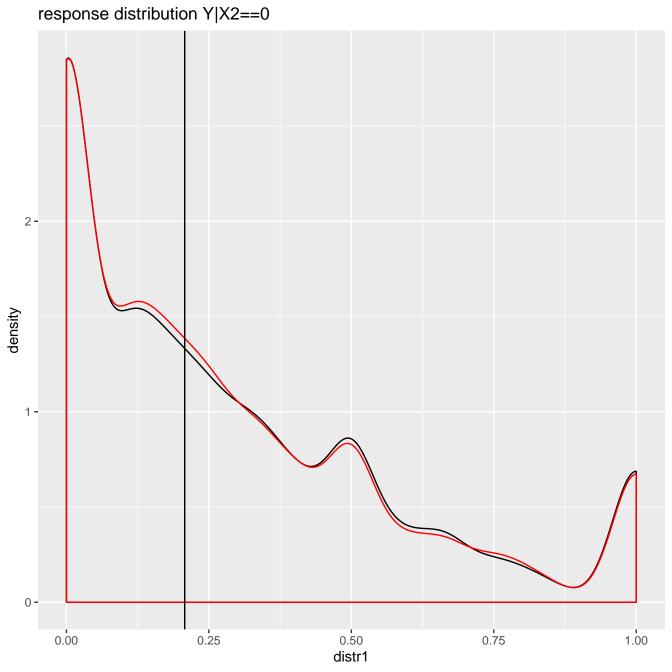


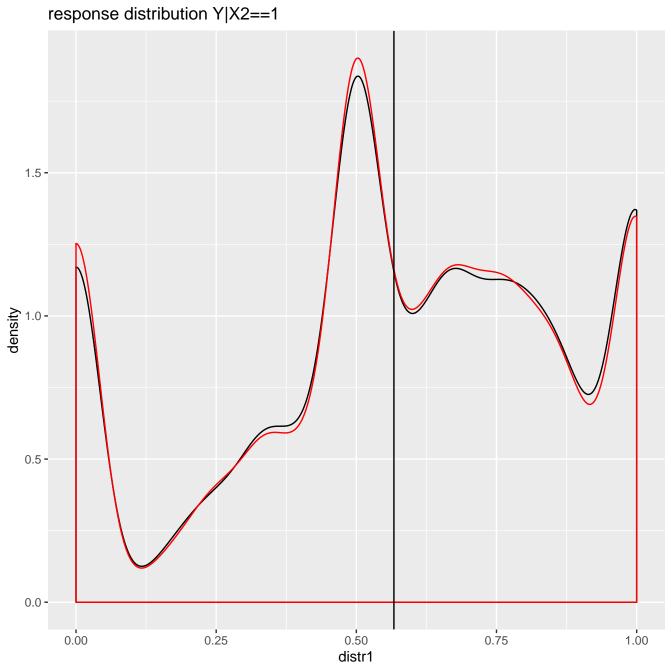


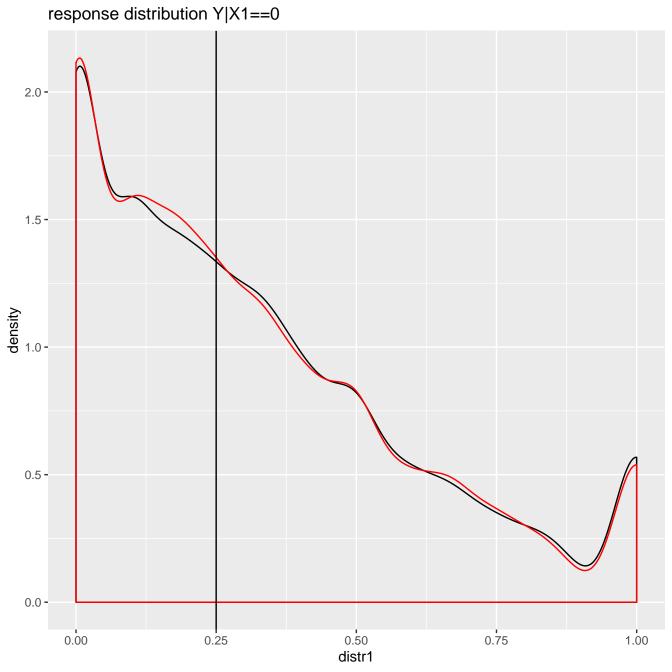


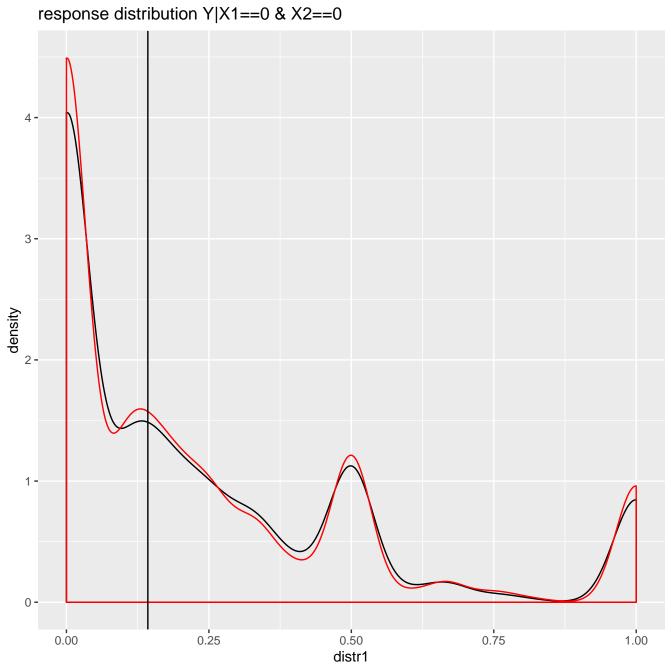
response distribution X1|Y==1 & X2==0 10.0 -7.5 density 5.0-2.5 -0.0 -0.25 0.50 **distr1** 0.75 0.00 1.00

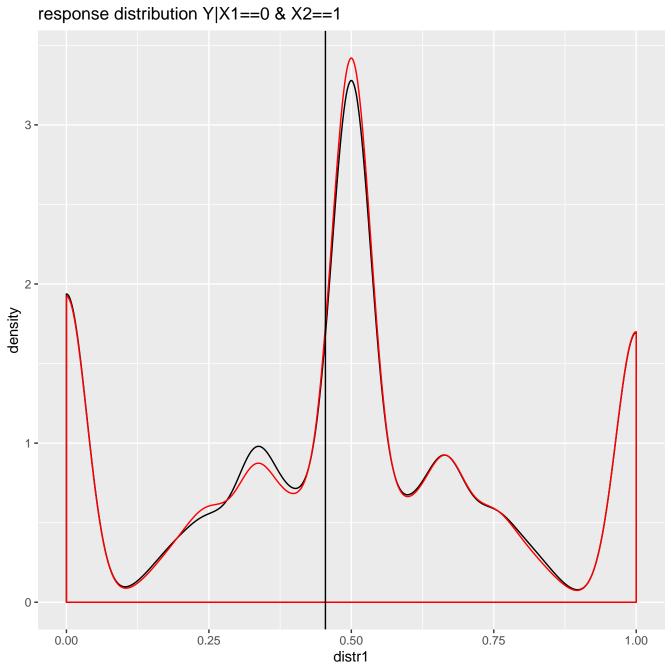


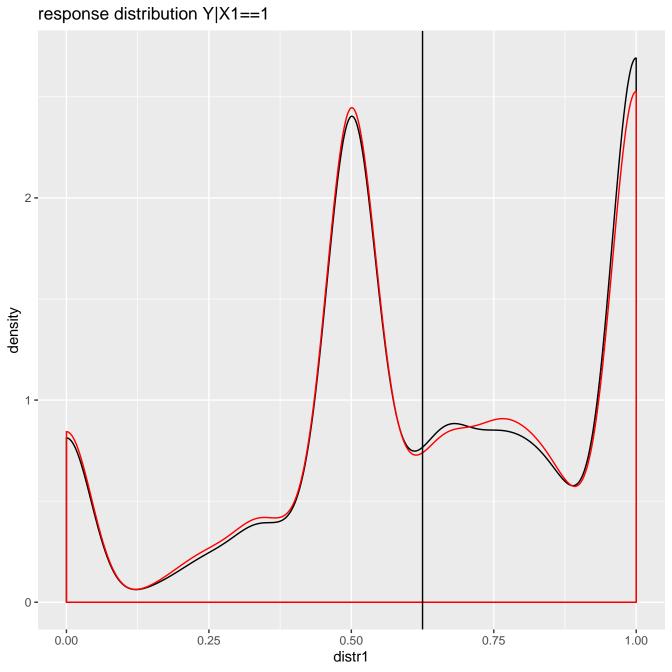


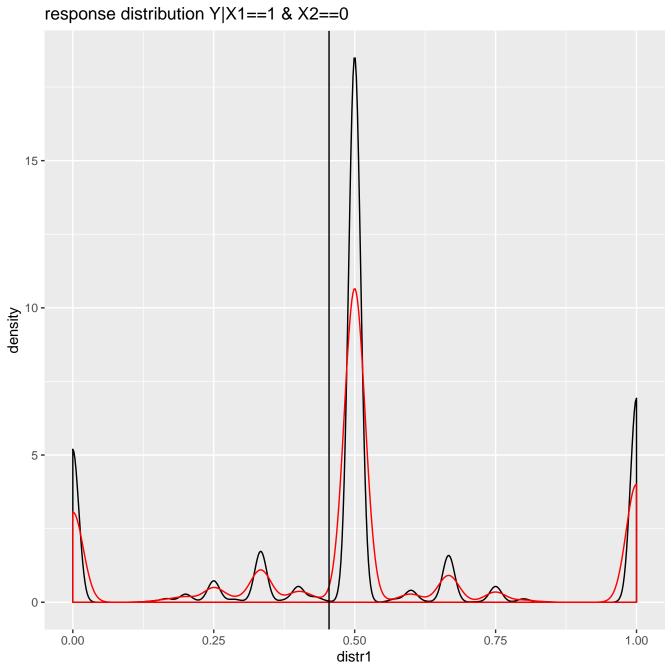


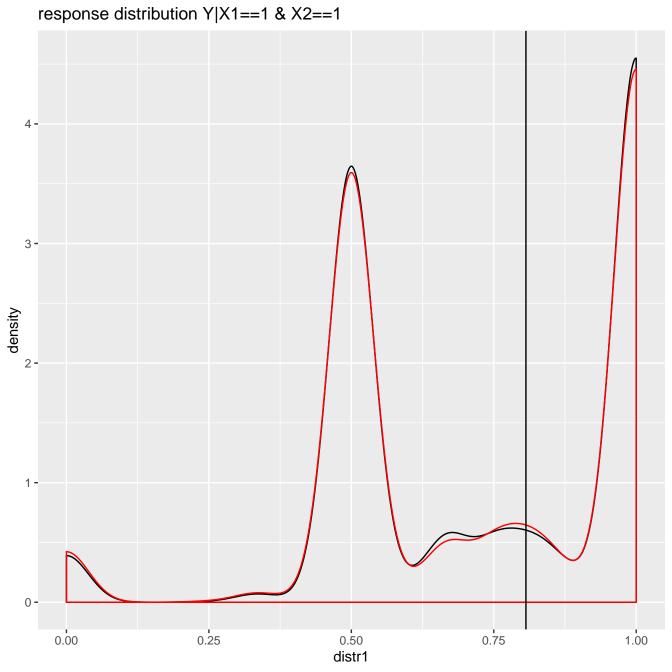




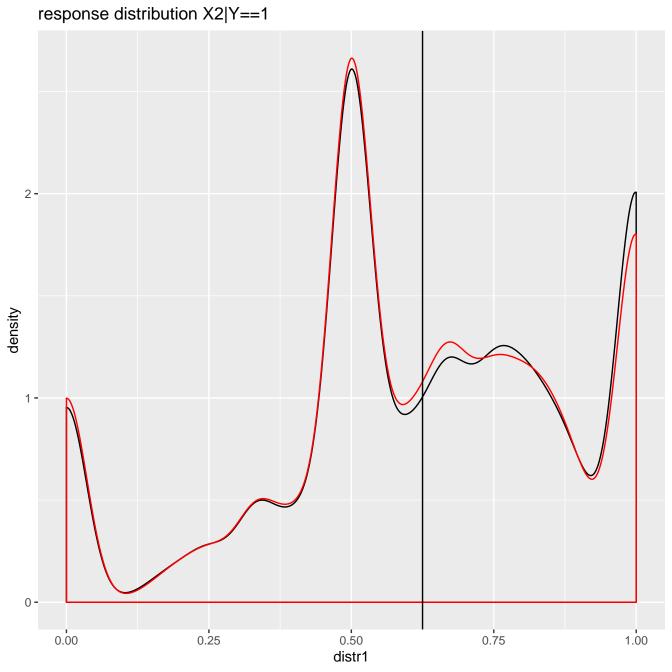


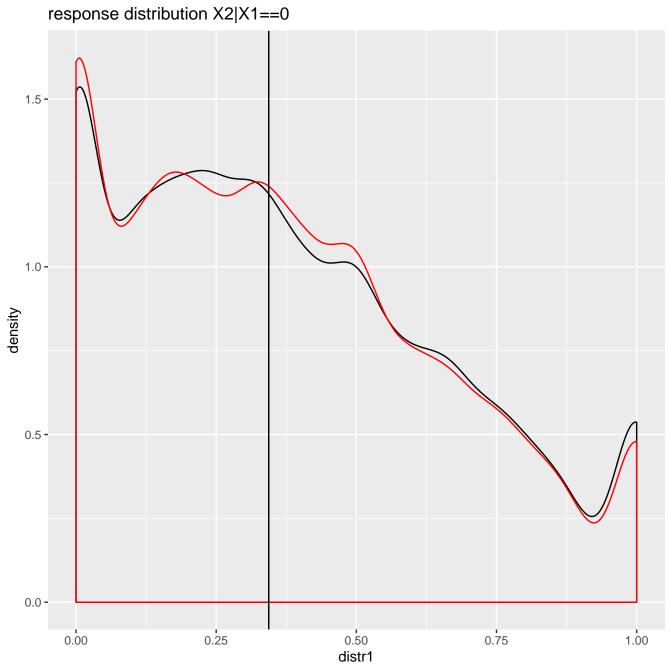






response distribution X2|Y==0 2.0 -1.5 density 1.0 -0.5 -0.0 -0.50 distr1 0.00 0.75 1.00 0.25





response distribution X2|Y==0 & X1==0 2.5 -2.0 -1.5 density 1.0 -0.5 -0.0 -0.50 1.00 0.00 0.75 0.25

distr1

