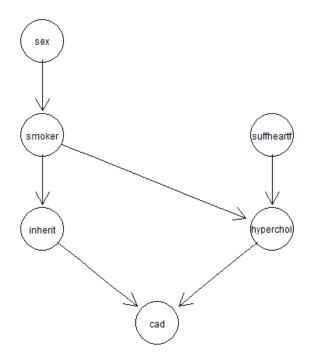
## Sai Kiran Putta

1)

a)

Building the network and plotting it gives the following plot.



b)
Joint probabilities before and after absorbing the evidence is given below.

```
> querygrain(grain1_compile, nodes = c("SuffHeartF", "CAD"), type = "joint")
    SuffHeartF
CAD     No     Yes
    No     0.3957368    0.1443930
    Yes    0.3118903    0.1479799
attr(,"class")
[1] "parray" "array"
> querygrain(grain1_compile_ev, nodes = c("SuffHeartF", "CAD"), type = "joint")
    SuffHeartF
CAD     No     Yes
    No     0.4078210    0.1453059
    Yes    0.2998061    0.1470670
attr(,"class")
```

Conditional probabilities before and after absorbing the evidence is given below.

```
> querygrain(grain1_compile, nodes = c("SuffHeartF", "CAD"), type = "conditional")
     SuffHeartF
CAD
             No
  No 0.7326698 0.2673302
  Yes 0.6782138 0.3217862
attr(,"class")
[1] "parray" "array"
> querygrain(grain1_compile_ev, nodes = c("SuffHeartF", "CAD"), type = "conditional")
     SuffHeartF
CAD
            No
  No 0.7373010 0.2626990
  Yes 0.6708976 0.3291024
attr(,"class")
[1] "parray" "array"
```

Marginal probabilities before and after absorbing the evidence is given below.

```
> querygrain(grain1_compile, nodes = c("SuffHeartF", "CAD"), type = "marginal")
$CAD
CAD
      No
                Yes
0.5401298 0.4598702
$SuffHeartF
SuffHeartF
      No
0.7076271 0.2923729
> querygrain(grain1_compile_ev, nodes = c("SuffHeartF", "CAD"), type = "marginal")
$CAD
CAD
               Yes
      No
0.5531269 0.4468731
$SuffHeartF
SuffHeartF
      No
0.7076271 0.2923729
```

Following are the 5 new data points that are generated.

```
> sim_find5
Sex Smoker Inherit CAD Hyperchol SuffHeartF

1 Female No No No No No
2 Female Yes Yes Yes Yes No
3 Female Yes Yes Yes Yes No
4 Female Yes Yes No No No
5 Female Yes No No Yes Yes
>
```

Following are the predictions of the newly generated 5 new data points.

```
$pred
$pred$Smoker
[1] "No" "Yes" "Yes" "Yes"

$pred$CAD
[1] "No" "Yes" "Yes" "No" "Yes"

$pEvidence
[1] 0.05544145 0.01984580 0.01984580 0.02075442 0.02882428
```

d) After generating new 500 data points and calculating the mis-classification rate, following are the results.

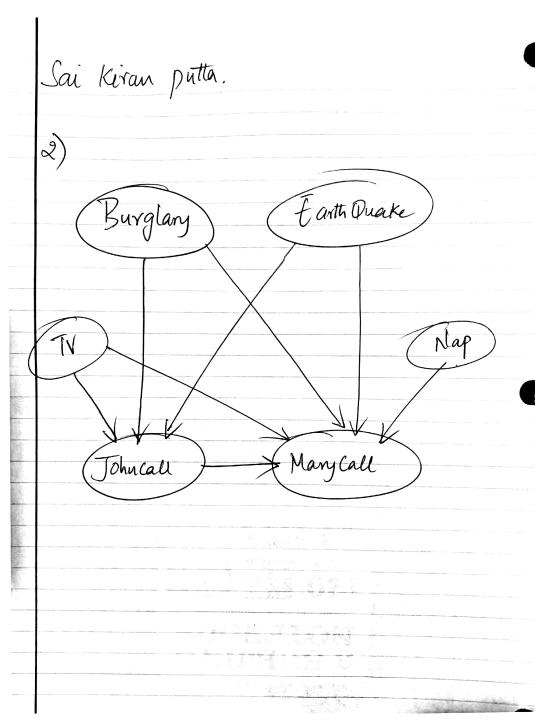
Mis-classification rate for smoker is 34.2 %

Mis-classification rate for CAD is 32.4 %

Since the performance of the network/classifier depends on the data, we cannot say if the accuracy is good enough. But the classifier doing  $\sim$ 15% better than a random guess.

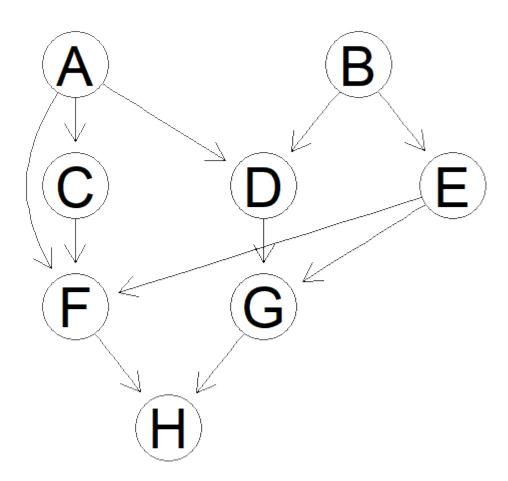
There might be other "valid combinations" of positions of nodes in the network which might lead to better accuracy.

Different versions could be tried to arrive at best performing network.



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Following is the network that is obtained.



Following are the results that were obtained for the questions.

- 'FALSE' A) C and G are d-separated.
- 'TRUE' B) C and E are d-separated.
- C) C and E are d-connected given 'FALSE' evidence about G.
- D) A and G are d-connected given 'TRUE'
- evidence about D and E.
- E) A and G are d-connected given 'FALSE' evidence on D.