
Advance Machine Learning CSE 674- Project 1

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Abstract

1 The goal of this project is to design probabilistic graphical models (PGMs) to
2 determine probabilities of observations which are described by several features. For
3 this project we took the handwriting patterns which are described by document
4 examiners. The PGM's can then be used to determine whether a particular hand-
5 writing sample is common (high probability) or rare (low probability) and which
6 in turn can be useful to determine whether a sample was written by a certain
7 individual. We considered only the letter pair **th** in this study since it is the most
8 commonly encountered pair of letters (called a bigram) in English. As a part of the
9 project we also evaluated **and** image dataset to construct a Bayesian network and
10 evaluate the goodness score (likelihood of a dataset) of several Bayesian networks
11 constructed from it.

12 **1 Feature definitions**

- 13 1) x1 - Height Relationship of t to h
- 14 2) x2 - Shape of Loop of h
- 15 3) x3 - Shape of Arch of h
- 16 4) x4 - Height of Cross on t staff
- 17 5) x5 - Baseline of h
- 18 6) x6 - Shape of t
- 19

20 **2 Evaluate pairwise correlations and independencies that exist in the data**

21 To determine if x_i and x_j are independent we can check if $P(x_i, x_j)$ is almost equivalent to $P(x_i) \times$
22 $P(x_j)$ where $P(x_i, x_j)$ is the joint probability distribution which is calculated as $P(x_i | x_j) \times P(x_j)$
23 . We can measure this closeness by approximately calculating the value using the below formula.

$$\sum abs((P(x, y) - P(x)P(y)))$$

24 We got the following values of closeness between the features to decide the independent sets that
25 might exist in the data.

- 26
- 27 1) $P(x_2 | x_1) = 0.15977$
- 28 2) $P(x_4 | x_1) = 0.11943$
- 29 3) $P(x_6 | x_1) = 0.16015$
- 30 4) $P(x_3 | x_2) = 0.218525$
- 31 5) $P(x_5 | x_2) = 0.132460$
- 32 6) $P(x_2 | x_3) = 0.2187580$

33 7) $P(x_5 | x_3) = 0.1155199$
34 8) $P(x_6 | x_3) = 0.095640$
35 9) $P(x_2 | x_4) = 0.119570$
36 10) $P(x_5 | x_4) = 0.115699$
37 11) $P(x_6 | x_4) = 0.1434699$
38 12) $P(x_2 | x_5) = 0.1312649$
39 13) $P(x_3 | x_5) = 0.1167000$
40 14) $P(x_1 | x_6) = 0.1603699$
41 15) $P(x_2 | x_6) = 0.1753150$
42 16) $P(x_3 | x_6) = 0.1390300$
43 17) $P(x_4 | x_6) = 0.1430700$

44

45 **From the above values we can get the independent sets by setting the significance level at**
46 **0.12. We consider the pairs whose closeness value is below 0.12 as the independent sets.**

47 **3 Construct various Bayesian network with the fewest number of edges that**
48 **maximizes the likelihood.**

49 3.1 Threshold

50 We set the threshold or significance level at **0.12** on the previous calculated result to determine if two
51 variables are independent .

52 3.2 Bayesian Model Creation

53 1)Based on the above results we generated 5 Bayesian models with the minimum edges possible and
54 the conditional probability distributions given to us for each node. To determine the goodness of the
55 models we used K2 score method provided by PGMPY library. The pre requisite to get the K2score is
56 to generate the sample data .

57

58 2)The sample data size of 5000 was generated where 1000 data size each was generated by
59 respective models and then combined and shuffled . This sample data is then used in K2score method
60 to determine the best model .

61

62 3)Ancestral sampling is used for data generation which is implemented by the class pgmpy.sampling.
63 BayesianModelSampling. This class is used to generate the data for the respective models.

64

65 3.2.1 Bayesian Network Model-1

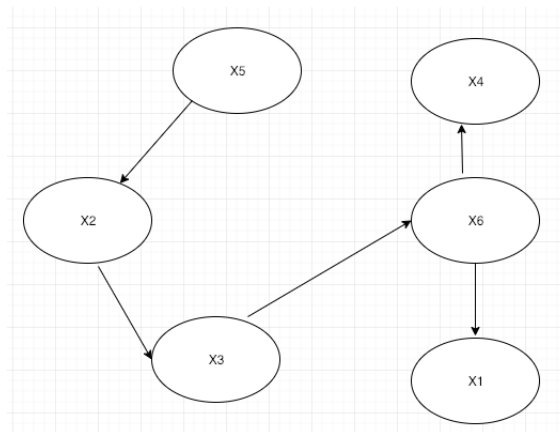


Figure 1: Bayesian Model 1

66 **3.2.2 Bayesian Network Model-2**

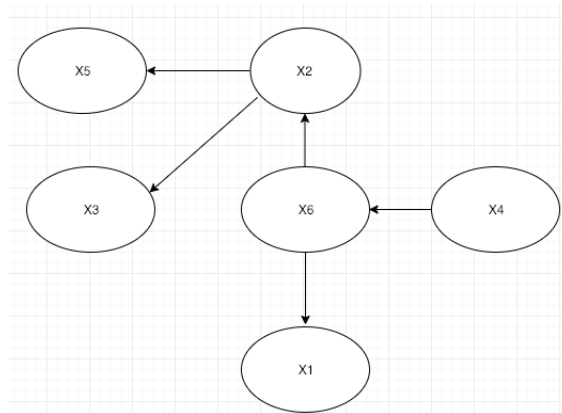


Figure 2: Bayesian Model 2

67 **3.2.3 Bayesian Network Model-3**

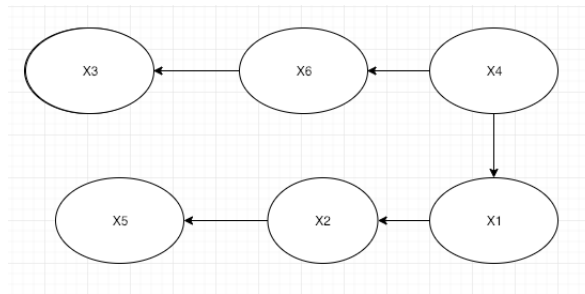


Figure 3: Bayesian Model 3

68 **3.2.4 Bayesian Network Model-4**

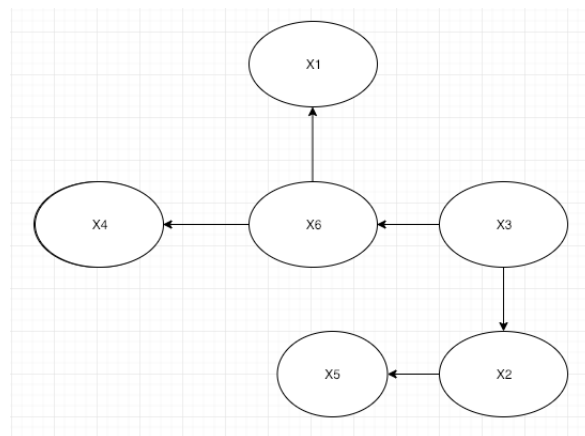


Figure 4: Bayesian Model 4

69 3.2.5 Bayesian Network Model-5

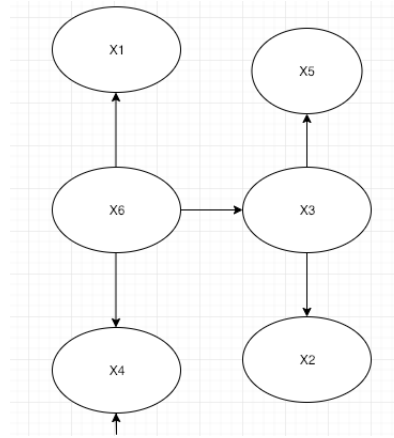


Figure 5: Bayesian Model 5

70 3.2.6 Model Performance Evaluation based on K2 Score

Bayesian Model 1 K2 Score	-31762.55432
Bayesian Model 2 K2 Score	-31785.13432
Bayesian Model 3 K2 Score	-32090.000980
Bayesian Model 4 K2 Score	-31761.52891
Bayesian Model 5 K2 Score	-31819.69284

71 Based on the above results we can say that Bayesian Model 4 performs the best among all of them.

72 3.2.7 Observations

73 1) High Probability TH :

X1	X2	X3	X4	X5	X6
0	1	1	0	3	3

74 2)Low Probability TH:

X1	X2	X3	X4	X5	X6
2	4	0	1	0	4

75 3) If there is a V structure being formed ,there was insufficient data to proceed for as for exam-
 76 ple $P(X_6 | P(X_1, X_4))$. We did not have the conditional probability distribution given 3 fea-
 77 tures.However as per the D separation we can rewrite $P(X_6 | P(X_1, X_4))$ as
 78 $P(X_1 | X_6) \times P(X_4 | X_6) \times P(X_6) | P(X_1, X_4)$. However it was observed that the conditional
 79 probability distribution did not sum up to 1 and was more than that,hence was not able to proceed
 80 with that particular Bayesian Network model.

81 4 Bayesian network into a Markov network using moralization.

82 1) Markov network is generated from the the best Bayesian Network model using the method
83 to_markov_model() given in the pgmpy library.

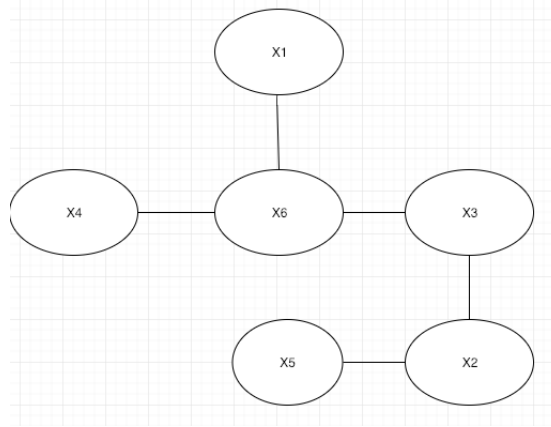


Figure 6: Markov Model

84 4.1 Inferences using Bayesian network and the Markov network

85 1) In inference we try to answer probability queries over the network given some other variables.
86 For example we might want to know the probable values of Shape of Arch of h given the Shape of
87 Loop of h. Variable Elimination library from pgmpy was used for inference for both the Bayesian and
88 the Markov network model. The example below depicts the inference process in both bayesian and
89 markov networks.

```
infer = VariableElimination(writerModel4)
print(infer.query(['x3'], evidence={'x5': 3}) ['x3'])
print(infer.query(['x2'], evidence={'x5': 3}) ['x2'])
print(infer.query(['x1'], evidence={'x5': 3}) ['x1'])
print(infer.query(['x4'], evidence={'x5': 3}) ['x4'])
print(infer.query(['x6'], evidence={'x5': 3}) ['x6'])
```

x3	phi(x3)
x3_0	0.1778
x3_1	0.6475
x3_2	0.1746

x2	phi(x2)
x2_0	0.2684
x2_1	0.3291
x2_2	0.0000
x2_3	0.1465
x2_4	0.2560

Figure 7: Inferences by Bayesian Model

```
infer_markov = VariableElimination(markov)
print(infer_markov.query(['x3'], evidence={'x5': 3}) ['x3'])
print(infer_markov.query(['x2'], evidence={'x5': 3}) ['x2'])
print(infer_markov.query(['x1'], evidence={'x5': 3}) ['x1'])
print(infer_markov.query(['x4'], evidence={'x5': 3}) ['x4'])
print(infer_markov.query(['x6'], evidence={'x5': 3}) ['x6'])
```

x3	phi(x3)
x3_0	0.1778
x3_1	0.6475
x3_2	0.1746
x2	phi(x2)
x2_0	0.2684
x2_1	0.3291
x2_2	0.0000
x2_3	0.1465
x2_4	0.2560

Figure 8: Inferences by Markov Model

1) Hence we can say that the inferences provided by the markov and bayesian models are very much accurate based on the data provided to us.

2) The computation time for bayesian and markov networks are quite similar but the markov network models are somewhat on the higher side than bayesian, although the difference is not much.

Bayesian Total Query Time for 100 queries in sec :1.05370900

Markov Total Query Time for 100 queries in sec :1.4753679

5 Construction of Bayesian and Markov networks based on the "and" image dataset

1) The data of 9 features of the and dataset was given to us.

2) To estimate the best model given the data ,we used HillClimbSearch technique that is available at from pgmpy.estimators.HillClimbSearch implements a greedy local search that starts from the DAG start (default: disconnected DAG) and proceeds by iteratively performing single-edge manipulations that maximally increases the score. The search terminates once a local maximum is found.Exhaustive Search is another technique to get the best model given data ,but it is usually not recommended when the number of feature is more than 5 which is 9 in our case;

3)The best model generated by HillClimbSearch along with the K2 score is as below:

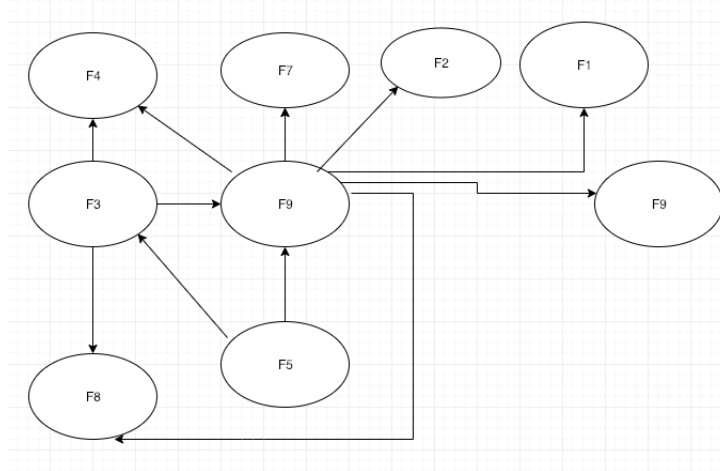


Figure 9: AND Features Bayesian Model by HillClimb Search

3) To try out different models , conditional independence is tested out using ConstraintBasedEstimator class and test_conditional_independence method between various features. We got the following results from it by keeping the significance level at 0.005 :

```
is_independent('f3', 'f4') - False
is_independent('f3', 'f9') - True
is_independent('f3', 'f8') - False
is_independent('f5', 'f9') - False
is_independent('f5', 'f3') - False
is_independent('f9', 'f8') - False
is_independent('f9', 'f7') - False
is_independent('f9', 'f1') - False
is_independent('f9', 'f6') - False
is_independent('f9', 'f2') - True
is_independent('f9', 'f4') - True
```

4) We constructed 4 more bayesian networks using the above independence relations:

Bayesian Model 1 :
('f3', 'f4'), ('f3', 'f8'), ('f5', 'f9'), ('f5', 'f3'), ('f9', 'f8'), ('f9', 'f7'), ('f9', 'f1'), ('f9', 'f6'), ('f9', 'f2')

Bayesian Model 2 :
('f3', 'f4'), ('f5', 'f9'), ('f5', 'f3'), ('f9', 'f8'), ('f9', 'f7'), ('f9', 'f1'), ('f9', 'f6')

Bayesian Model 3 :
('f4', 'f3'), ('f9', 'f5'), ('f3', 'f5'), ('f8', 'f9'), ('f7', 'f9'), ('f1', 'f9'), ('f6', 'f9')

Bayesian Model 4 :
('f4', 'f3'), ('f3', 'f8'), ('f5', 'f9'), ('f5', 'f3'), ('f9', 'f8'), ('f9', 'f7'), ('f9', 'f1'), ('f9', 'f6'), ('f2', 'f9')

140 5.1 K2 Score Evaluation

AND Model	K2 Score
Hill Climb Search Model	-9462.7048
Bayesian Model 1	-9472.1900
Bayesian Model 2	-8170.1712
Bayesian Model 3	-8332.6588
Bayesian Model 4	-9504.2481

141 5.2 Observations

142 1) If we consider that (f2 , f9) and (f4 , f9) features are independent based on the significance level
143 the k2 score improves to -8170.1772.

144

145 2)But we cannot conclude this as the data might be less to determine the correlation be-
146 tween the features.

147

148 3)Hence we consider Hill Climb Search model as the best model among them.

149

150 5.3 Markov Model Generation

151 1) Markov model is generated from the best bayesian model using moralization method
152 to_markov_model() from the pgmpy library.

153 5.4 Conditional Probability Generation

154 1) We generated the condition probability distributions for the best model using MaximumLikeli-
155 hoodEstimator and BayesianEstimator. Slight difference was observed between the 2 methods as
156 BayesianEstimator uses K2 prior, which simply adds 1 to the count of every single state and then
157 later updated using the state counts from the observed data.

158

159 2)The approach in Maximum Likelihood Estimation (MLE) is to fill the CPDs in such a
160 way, that $P(data \mid model)$ is maximal and uses relative frequencies.

161

162 5.5 Inference

163 1) Variable Elimination method is used to generate inference given certain evidence of the fea-
164 tures.Similar results was observed for both bayesian and markov model.

```
from pgmpy.inference import VariableElimination
infer = VariableElimination(best_model)
print(infer.query(['f1'], evidence={'f9': 0}) ['f1'])
```

f1	phi(f1)
f1_0	0.1991
f1_1	0.4459
f1_2	0.2641
f1_3	0.0909

Figure 10: Inferences example for the **and** model

165 5.6 Likelihood of the dataset

166 1)For both the markov and bayesian models ,we can verify that the inference produced by the models
167 matches with the estimated conditional probability distributions as shown below.

f9	f9(1)	f9(2)
f1(0)	0.19913419913419914	0.10709838107098381
f1(1)	0.4458874458874459	0.2602739726027397
f1(2)	0.26406926406926406	0.3686176836861768
f1(3)	0.09090909090909091	0.2640099626400996

Figure 11: CPD for $P(F1 | F9)$

f1	phi(f1)
f1_0	0.1991
f1_1	0.4459
f1_2	0.2641
f1_3	0.0909

Figure 12: Inferences query for F1 given F9 as 0

168 2)Hence we can conclude that dataset of "and" fetaures is quite relevant and produces inference as
169 per the expectation.
170

171 3)The computation time for both the bayesian and markov model is also very similar.
172

173 4) The computation time for both the bayesian and markov models are similar with the markov model
174 being slightly on the higher side.

175 Bayesian And Model Total Query Time for 100 queries in sec :2.2328779999999

176 Markov And Model Total Query Time for 100 queries in sec :2.5165949999999999

177

178 References

179 [1] <http://pgmpy.org/>