ExpertBayes: Automatically Refining Manually Built Bayesian Networks



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- Objectives
- Dataset
- Methodology and Tools
- Results and Analysis
- *ExpertBayes* (graphical user interface)
- Conclusions and Future Work



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Dataset

- Prostate Cancer:
 - ≻496 cases

> Each case refers to the clinical history of each patient

- Breast Cancer (1) :
 - ≻100 cases

Each case refers to a breast nodule from mammography results

- Breast Cancer (2) :
 - ►241 cases

Each case refers to a breast nodule from mammography results

Attributes

Prostate Cancer



11 Attributes

Age (age)

Weight (wt)

Family history of cancer (hx)

Systolic blood pressure (Sbp)

Diastolic blood pressure (Dbp)

Hmoglobins (hg)

Clinical stage (stage)

Doubling time PSA (Dtime)

Size of the prostate (size)

Bony metastases (bm)



145 Alive

(-)

351 Dead (+)

Attributes





Age

Disease

BreastDensity

MassesShape

MassesDensity

MassesSize

PostOpChange

MassesStability

Calc_Milk



Attributes





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Methodology and Tools





- 5-fold cross-validation to train and test our models
- **t-test** was used to validate the results
 - Significance level: 0.05



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• CCI(%) test set - averaged across 5-folds

Dataset	Original	ExpertBayes	WEKA-K2	WEKA-TAN
Prostate Cancer	74	76	74	71
Breast Cancer (1)	49	63	59	57
Breast Cancer (2)	49	64	80	79



- Precision-Recall Curves for various thresholds
 - Prostate





- Precision-Recall Curves for various thresholds
 - Breast Cancer (1)





- Precision-Recall Curves for various thresholds
 - Breast Cancer (2)







CCI :74%

CCI :76%



Weka TAN





CCI :71%

CCI :76%



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ExpertBayes

- Allow the user :
 - Load new Network;
 - Load new data;
 - Load new tables of conditional probabilities;
 - Save the network;
 - Add / Remove vertex;
 - Add / Remove edge;
 - Return edge;
 - Visualize the score, confusion matrix, CPT of an node, precision-recall curve and ROC curve;
- Graphical user interface



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Conclusions and Future Work

- ExpertBayes produces better results than the original model and better results than models learned with other tools.
- ExpertBayes also provides a graphical user interface (GUI) where users can play with their models thus exploring new structures that give rise to a search for other models.



Conclusions and Future Work

- Improve the algorithm in order to have better prediction performance.
- Using more (and quality) data, different search and parameter learning methods.

Thank you!





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Appendices



State of the Art

- Previous works considered as initial network a naive Bayes or empty network [9], [4]:
 - [9] Hall, M., Frank, E., Holmes, G., Pfahringer, B., Reutemann, P., Witten,
 I.H.: The weka data mining software: an update. SIGKDD Explor. Newsl. 11, 10–18 (Nov. 2009), 1656274.1656278
 - [4] Chan, H., Darwiche, A.: Sensitivity analysis in bayesian networks: From single to multiple parameters. In: Proceedings of the 20th Conference on Uncertainty in Artificial Intelligence. pp. 67–75. UAI '04, AUAI Press, Arlington, Virginia, United States (2004),id=1036843.1036852



State of the Art

- The R packages deal [2] and bnlearn [11], [13] can refine any input network. However, deal and bnlearn refine input networks by successive refinements instead of performing the refinement only over the original network:
 - [2] Bottcher, S.G., Dethlefsen, C.: Deal: A package for learning bayesian networks. Journal of Statistical Software 8, 200–3 (2003)
 - [11] Nagarajan, R., Scutari, M., Lebre, S.: Bayesian Networks in R with Applications in Systems Biology. Springer, New York (2013), iSBN 978-1461464457
 - [13] Scutari, M.: Learning bayesian networks with the bnlearn R package. Journal of Statistical Software 35(3), 1–22 (2010), http://www.jstatsoft.org/v35/i03/



State of the Art

- WEKA, whose bayesian algorithms apply successive refinements to the newly built models:
 - [6] Cooper, G.F., Herskovits, E.: A bayesian method for the induction of probabilistic networks from data. Machine Learning 9(4), 309–347 (1992), BF00994110
 - [8] Friedman, N., Geiger, D., Goldszmidt, M.: Bayesian network classifiers.
 In: Machine Learning. vol. 29, pp. 131–163 (1997)



Methodology

WEKA :

• K2 is a greedy algorithm that, given an upper bound to the number of parents for a node, tries to find a set of parents that maximizes the likelihood of the class variable [6].

• TAN (Tree Augmented Naive Bayes) generates a tree over naive Bayes structure, where each node has at most two parents, being one of them the class variable [8].



Data Distribution

Dataset	Number of Instances	Number of Variables	Pos.	Neg.
Prostate Cancer	496	11	352	144
Breast Cancer (1)	100	34	55	45
Breast Cancer (2)	241	8	88	153

CRACS

The pseudo-code for ExpertBayes

Data:

OriginalNet, // initial network structure; Train // training set;

Test // test set

Result:

scoreTrain // scores in the training set for BestNet
scoreTest // scores in the test set for BestNet
BestNet // best scored network on Train

- 1 Read OriginalNet;
- 2 Read Train and Test sets;
- 3 BestNet = OriginalNet;
- 4 Learn parameters for OriginalNet from training set;

5 repeat

- 6 Randomly choose a pair of nodes N_1 and N_2 ;
- 7 **if** there exists an edge between N_1 and N_2 then
- 8 randomly choose: revert or remove
- 9 else
- 10 choose add operation;
- 11 randomly choose edge direction

12 end

- 13 Apply operation to OriginalNet obtaining NewNet;
- 14 Rebuild necessary CPT entries, if necessary;
- 15 Compute scoreTrain of the NewNet;
- 16 if scoreTrain NewNet > scoreTrain BestNet then
- 17 BestNet = NewNet
- 18 end
- 19 until N iterations using OriginalNet and Train;
- 20 Apply BestNet to Test and compute scoreTest;



ExpertBayes Advantages

- Reduces the computational costs;
- Embed knowledge of an expert in the newly built network;
- Allows the construction of fresh new networks, through its graphical interface.