



CIS 830: Advanced Topics in Artificial Intelligence













Algorithm Max-Spanning-Tree-Structure - Estimate P(x) and P(x, y) for all single random variables and pairs; |(X; Y) = D_{xL}(P(X, Y) || P(X) · P(Y)) Build complete undirected graph: variables as vertices, I(X; Y) as edge weights - T ← Build-MWST (V × V, Weights) // Chow-Liu algorithm: weight function = I - Set directional flow on T and place the CPTs on its edges (gradient learning) - RETURN: tree-structured BBN with CPT values - Advantage: Restricts hypothesis space and limits overfitting capability - Disadvantage: It only searches a single parent and some available data may be lost The "Sparse Candidate" Algorithm - It builds a network structure with maximal score by limiting H to at most K parents for each variables in BBN (K < N)</td> - Searching Candidate sets K: Based on D and B_{n+1}, select for each variables X₁ as et of



Maximize : Find a network 9, maximizing score (9, 10) among networks
 Advantages: Overcoming the drawbacks of MSTS algorithm
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 Description

Learning Structure





	Learning with Small Candidate Sets
•	Maximal Restrict Bayesian Network (MRBN)
	 Input: A set D = {X¹,, Xⁿ} of instances; a digraph H of bounded in-degree K; and a decomposable score S
	– Output: A network B = <g, <math="" display="inline">\ominus> so that G \subseteq H, that maximizes S with respect to D</g,>
•	Standard Heuristics
	 No knowledge of expected structure, local change (e.g. arc deletion, arc addition, and arc reversal), and local maximum score
	 Algorithms: Greedy hill-climbing; Best-first search; and Simulated annealing
	 Time complexity In Greedy hill climbing is O(n²) for initial change, then becomes linear O(n) for each iteration
	- Time complexity in MRBN is O(kn) for initial calculation, then becomes O(k)
•	Divide and Conquer Heuristics
	- Input: A digraph H = {X _i -> X _i : X _i \in C _i }, and a set of weights w(X _i , Y) for each X _i , Y \in C _i
	- Output: An acyclic subgraph $G \subseteq H$ that maximizes $W_H[G] = \sum_i w(X_i, Pa(X_i))$
	 Decompose H by using standard graph decomposition methods
	- Find a local maximum weight
	- Combine them into a global solution.
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Decomposition

Strongly Connected Components: (SCC)

- A subset of vertices A is strongly connected if for each X, $Y \in A$, there is a directed path from X to Y and a directed path from Y to X
- Decomposition of SCC into maximal sets that have no strongly connected components

Separator Decomposition

- Searching a separator of H which separate H into H1 and H2 with no edges between them
- **Cluster-Tree Decomposition**
- Cluster tree definition
- Decomposing into cluster tree
- **Cluster-Tree Heuristic**
- A mixture of cluster-tree decomposition algorithm and standard heuristics
- Using for the decomposition of H for large size clusters KSI

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Experimental Evaluation

 Using TABU search to find 	Method	Iter	Time	Score	KL	Stats
global max score	Greedy		40	-15.35	0.0499	2656
 "Alarm" network 	Disc 5	1	14	-18.41	3.0608	908
– Samples: 10000		2	19	-16.71	1.3634	1063
– variables: 37		3	23	-16.21	0.8704	1183
- including 13 have 2 values, 22	Disc 10	1	20	-15.53	0.2398	1235
have 3 values, and 2 have 4		2	26	-15.43	0.1481	1512
values		3	32	-15.43	0.1481	1733
Text Test	Shld 5	1	14	-17.50	2.1675	915
– Samples: 20 * 1000 sets		2	29	-17.25	1.8905	1728
12		3	36	-16.92	1.5632	1907
5.	shid 10	1	20	-15.86	0.5357	1244
ffreener /		2	35	-15.50	0.1989	1968
		3	41	-15.50	0.1974	2109
3 <i>{</i>	Score 5	1	12	-15.94	0.6756	893
Greedy HC		2	27	-15.34	0.0550	1838
5 Disc 5 Disc 15		3	34	-15.33	0.0479	2206
Score 5 Score 15 - 0 Shid 5	Score 10	1	17	-15.54	0.2559	1169
4 1. 11 / Shid 15		2	30	-15.31	0.0352	1917
0 200 400 600 800 1000 1200 1400 1600 1800 2000 Time		3	34	-15.31	0.0352	2058
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Summary

Content Critique

- Key Contribution It presents an algorithm to select candidate sets and to discover efficiently the maximum score of Bayesian networks.
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- efficiently the maximum score of Bayesian networks. Strengths
 I tuses scoring measure instead of mutual information to measure the dependency of parent and children, then uses the maximum score to build BBN This algorithm can allow children to have multiple parents and handle random variables with multiple values. The limited candidate sets provide a small hypothesis space The time complexity of searching the maximum score in BBN is linear I tis sepacially efficient for massive datasets Weaknesses I denerif complete the optimizer of sources of component of complexity of searching the maximum score in BBN is linear

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 - It doesn't consider the existing of spurious dependency of random variables
 The search of candidate sets is complex.
 It is no better for small datasets than standard heuristic algorithms

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Presentation Critique

- Selfation Critique
 Audiences: Medical diagnosis; Mapping learning; language understanding; Image processing
 Positive points: Presents a useful approach in building BBN structure
 Negative points: No comparison with other algorithms

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