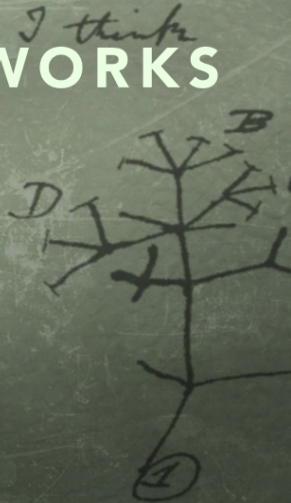
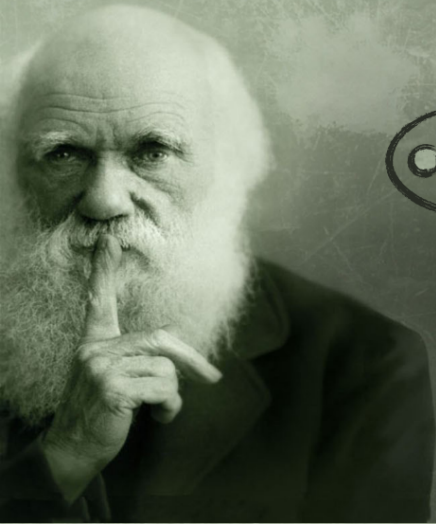


INTRODUCING

# DARWINIAN NETWORKS



*There between*

# Introducing Darwinian Networks

Cory J. Butz   Jhonatan S. Oliveira   André E. dos Santos

Department of Computer Science  
University of Regina  
Regina, S4S 0A2, Canada

[www.darwiniannetworks.com](http://www.darwiniannetworks.com)

28<sup>th</sup> FLAIRS Conference  
Hollywood, Florida, USA  
May 18<sup>th</sup>, 2015

# Outline of the Presentation

1. Motivation
2. Darwinian Networks
  - 2.1 Inference
    - Variable Elimination
    - Arc-Reversal
    - Lazy Propagation
  - 2.2 Modeling (Testing Independencies)
    - m-Separation
    - d-Separation
3. Advantages
4. Conclusions

# 1. Motivation

Understanding *Bayesian network* (BN) inference is not easy.

We sought a *purely graphical approach* for BN inference.

The representation took on a biological feel (*Darwinian Networks*).

We then observed that Darwinian Networks could represent, simplify, and speed-up the testing of independencies.

And determine good elimination orderings.

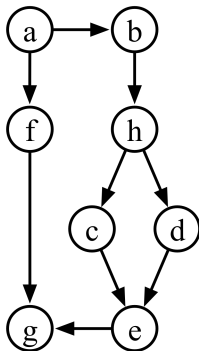
Surprisingly simple, remarkably robust.

# Bayesian Networks

A *Bayesian Network* (BN) consists of:

- a *directed acyclic graph* (DAG),
- a matching set of *conditional probability tables* (CPTs).

## Example: BN



$$P(U) = P(a) \cdot P(b|a) \cdot P(c|h) \cdot P(d|h) \cdot P(e|c, d) \cdots P(g|e, f)$$

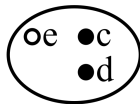
## 2. Darwinian Networks (DNs)

A CPT  $P(X|Y)$  is represented as a *population*  $p(X, Y)$ .

The variables in the LHS  $X$  are *white*.

The variables in the RHS  $Y$  are *black*.

$$P(e|c, d)$$

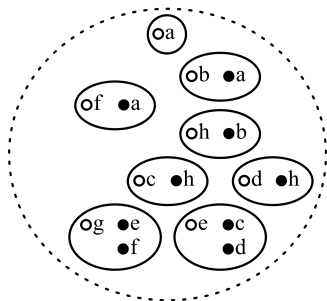


CPT  $P(e|c, d)$  is depicted as population  $p(e, cd)$ .

## 2. Darwinian Networks

A *Darwinian Network* (DN) is a finite, multiset of populations.

A DN is depicted by a dashed closed curve around its populations.

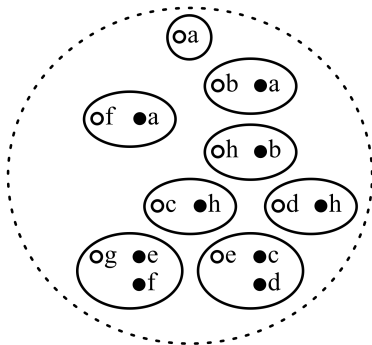
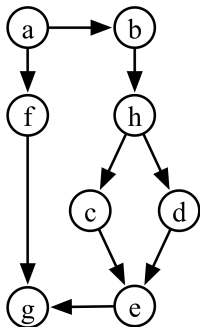


Populations:

$p(a)$ ,  $p(b, a)$ ,  $p(c, h)$ ,  $p(d, h)$ ,  $p(e, cd)$ ,  $p(f, a)$ ,  $p(h, b)$ ,  $p(g, ef)$



# Every BN can be represented as a DN



## 2.1 Inference

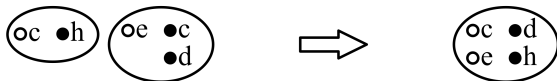
BN inference is called *evolution* in DNs.

We first introduce operations on populations corresponding to:

- multiplication,
- division,
- marginalization.

# Multiplication is the Merge of Populations

$$P(c|h) \cdot P(e|c, d) = P(c, e|d, h)$$



○ white + ● black = ○ white

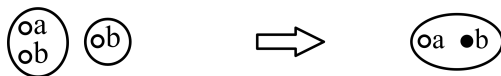
● black + ○ white = ○ white

● black + ● black = ● black

○ white + ○ white = ● black

## Merge also represents Division

$$P(a, b) / P(b) = P(a|b)$$



○ white + ● black = ○ white

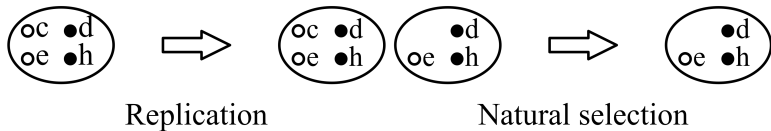
● black + ○ white = ○ white

● black + ● black = ● black

○ white + ○ white = ● black

# Marginalization is Replication then Natural Selection

$$\sum_c P(c, e|d, h) = P(e|d, h)$$



# DNs can represent BN Inference Algorithms

As DNs can represent multiplication, division, and marginalization, it follows that DNs can represent exact inference algorithms, including:

- Variable Elimination (VE) (Zhang and Poole, 1994),
- Arc-Reversal (AR) (Olmsted, 1983),
- Lazy Propagation (LP) (Madsen and Jensen, 1999).

# Variable Elimination (VE)

To answer  $P(e|b)$ , VE computes:

$$P(c, e|d, h) = P(c|h) \cdot P(e|c, d), \quad (1)$$

$$P(e|d, h) = \sum_c P(c, e|d, h), \quad (2)$$

$$P(d, e|h) = P(d|h) \cdot P(e|d, h), \quad (3)$$

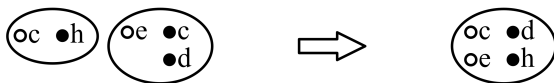
$$P(e|h) = \sum_d P(d, e|h), \quad (4)$$

$$P(e, h|b) = P(h|b) \cdot P(e|h), \quad (5)$$

$$P(e|b) = \sum_h P(e, h|b). \quad (6)$$

## Example: Representing VE as DN Evolution

$$P(c|h) \cdot P(e|c, d) = P(c, e|d, h) \quad (1)$$

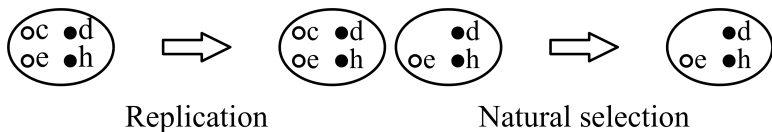


Merge



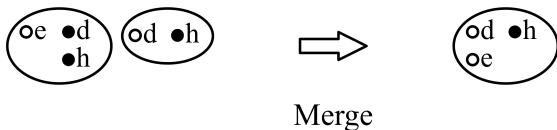
## Example: Representing VE as DN Evolution

$$\sum_c P(c, e|d, h) = P(e|d, h) \quad (2)$$



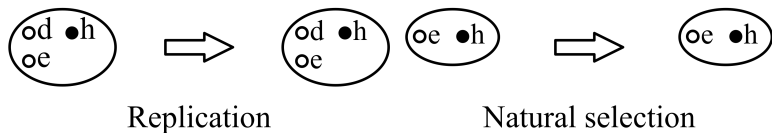
## Example: Representing VE as DN Evolution

$$P(e|d, h) \cdot P(d|h) = P(d, e|h) \quad (3)$$



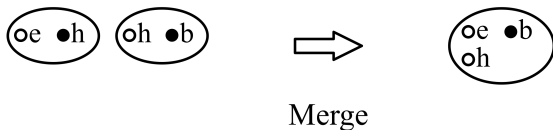
## Example: Representing VE as DN Evolution

$$\sum_d P(d, e|h) = P(e|h) \quad (4)$$



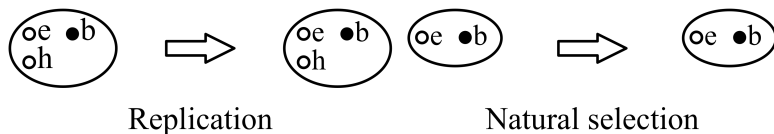
## Example: Representing VE as DN Evolution

$$P(e|h) \cdot P(h|b) = P(e, h|b) \quad (5)$$



## Example: Representing VE as DN Evolution

$$\sum_h P(e, h|b) = P(e|b) \quad (6)$$



Query  $P(e|b)$  is represented as population  $p(e, b)$ .

# DN Advantage

Koller and Friedman (2009) introduce readers to BN inference using VE.

There is a **one-to-one correspondence** between VE's mathematical equations and the DN illustrations.

Hence, one can learn VE without involving a single equation.

## DNs can represent Arc-Reversal (AR)

AR eliminates a variable  $v_i$  by reversing the arc  $(v_i, v_j)$  between  $v_i$  and each child  $v_j$  of  $v_i$ .

$$\begin{aligned}P(v_i, v_j | P_i P_j) &= P(v_i | P_i) \cdot P(v_j | P_j), \\P(v_j | P_i P_j) &= \sum_{v_i} P(v_i, v_j | P_i P_j), \\P(v_i | P_i P_j v_j) &= \frac{P(v_i, v_j | P_i P_j)}{P(v_j | P_i P_j)}.\end{aligned}$$

AR only involves multiplication, division, and marginalization.

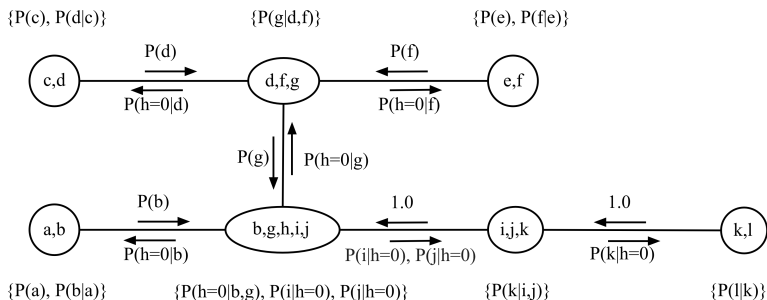
# Lazy Propagation (LP)

BN variables are clustered into nodes, organized as a join tree.

Each BN CPT is assigned to a join tree node.

Messages are propagated systematically.

LP only involves multiplication, division, and marginalization.



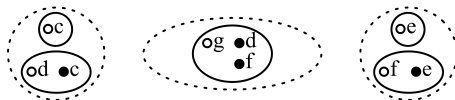


## DNs can represent LP

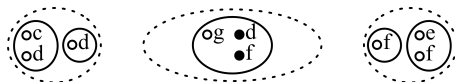
A join tree is represented as a set of DNs.

Each join tree node is represented as one DN.

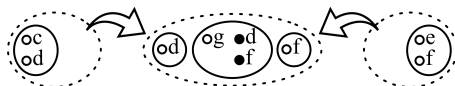
A propagated message from a join tree node to another is viewed as a population *migrating* from one DN to another.



Evolution



Migration

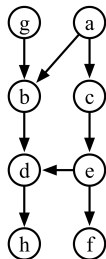


## 2.2 Modeling - Testing Independencies in BNs

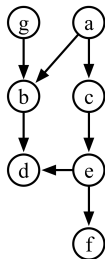
*m-Separation* (Lauritzen et al., 1990; Zhang and Poole, 1994) tests  $I(X, Y, Z)$  in an undirected graph with four steps:

- (i) construct the sub-DAG onto  $XYZ \cup An(XYZ)$ ;
- (ii) construct the *moralization* by adding an undirected edge between each pair of parents of a common child and then dropping directionality;
- (iii) delete  $Y$  and its incident edges;
- (iv) if there exists a path from  $X$  to  $Z$ , then  $I(X, Y, Z)$  does not hold; otherwise,  $I(X, Y, Z)$  holds.

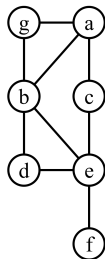
# Example: m-Separation $I(a, d, f)$



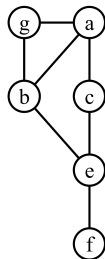
DAG



sub-DAG



moralization



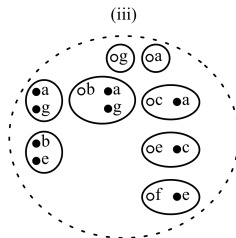
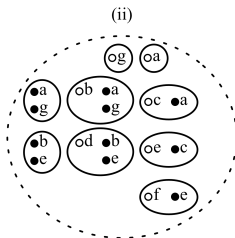
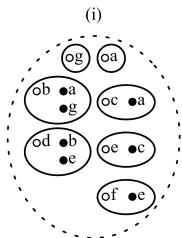
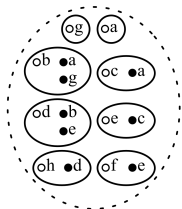
deletion  
and  
check for path

# DNs can represent m-Separation

Testing independencies in BNs is represented as testing *adaptation* in DNs.

We observe how populations *adapt* to the removal of other populations.

# Adaptation: Testing Independence $I(a, d, f)$ in DNs

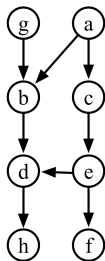


## DNs Simplify m-Separation

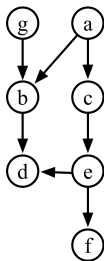
Moralization of m-separation can be excessive.

Adding edge  $(b, e)$  is necessary, since  $d$  will be deleted.

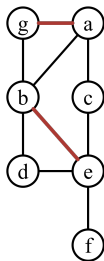
Adding edge  $(a, g)$  is unnecessary, since  $b$  will not be deleted.



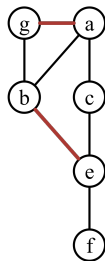
DAG



sub-DAG



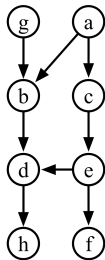
moralization



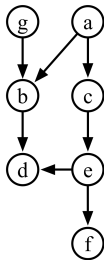
deletion  
and  
check for path

# DN Contribution (Rationalization)

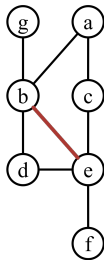
When testing  $I(X, Y, Z)$ , add an undirected edge between variables with a common child **only when the child is in  $Y$** .



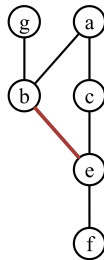
DAG



sub-DAG



rationalization



deletion  
and  
check for path

# Modeling - Testing Independencies with d-Separation

**Algorithm 3.1** Find nodes reachable from  $X$  given  $Y$  via active paths in DAG  $\mathcal{B}$

```
1: procedure REACHABLE( $X, Y, \mathcal{B}$ )
2:    $\triangleright$  Phase I: insert  $Y$  and all ancestors of  $Y$  into  $A$ 
3:    $An(Y) \leftarrow ANCESTORS(Y, \mathcal{B})$ 
4:    $A \leftarrow An(Y) \cup Y$ 
5:    $\triangleright$  Phase II: traverse active paths starting from  $X$ 
6:   for  $v \in X$  do  $\triangleright$  (Node, direction) to be visited
7:      $L \leftarrow L \cup \{(\uparrow, v)\}$ 
8:    $V \leftarrow \emptyset$   $\triangleright$  (Node, direction) marked as visited
9:    $R \leftarrow \emptyset$   $\triangleright$  Nodes reachable via active path
10:  while  $L \neq \emptyset$  do  $\triangleright$  While variables to be checked
11:    Select  $(d, v)$  in  $L$ 
12:     $L \leftarrow L - \{(d, v)\}$ 
13:    if  $(d, v) \notin V$  then
14:      if  $v \notin Y$  then
15:         $R \leftarrow R \cup \{v\}$   $\triangleright v$  is reachable
16:       $V \leftarrow V \cup \{(d, v)\}$   $\triangleright$  Mark  $(d, v)$  as visited
17:      if  $d = \uparrow$  and  $v \notin Y$  then
18:        for  $v_i \in Pa(v)$  do
19:           $L \leftarrow L \cup \{(\uparrow, v_i)\}$ 
20:        for  $v_i \in Ch(v)$  do
21:           $L \leftarrow L \cup \{(\downarrow, v_i)\}$ 
22:      else if  $d = \downarrow$  then
23:        if  $v \notin Y$  then
24:          for  $v_i \in Ch(v)$  do
25:             $L \leftarrow L \cup \{(\downarrow, v_i)\}$ 
26:        if  $v \in A$  then
27:          for  $v_i \in Pa(v)$  do
28:             $L \leftarrow L \cup \{(\uparrow, v_i)\}$ 
29:  return  $R$ 
```

Geiger et al. (1989) provide a linear time complexity algorithm for implementing d-separation.

Implementation of d-separation given by Koller and Friedman (2009).



## DNs Simplify d-Separation

The main idea is to start from  $X$ , follow active paths, and see if it reaches  $Z$ .

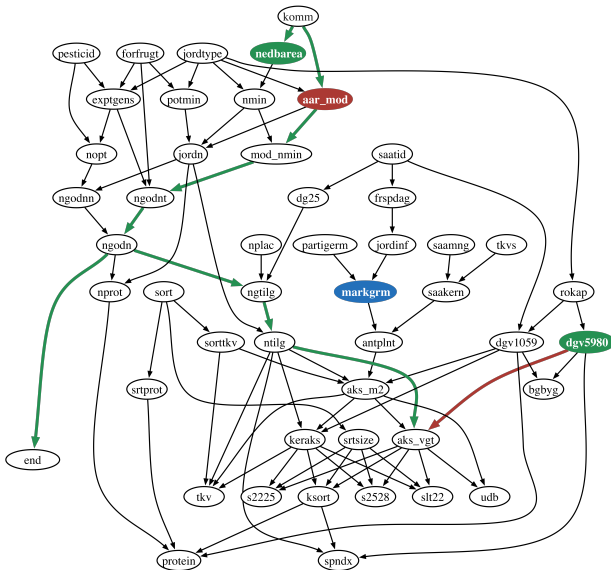
The linear implementation of d-separation considers all active paths until they become blocked.

The DN improvement is the identification of a class of active paths that are **doomed to become blocked**.

Thus, there is no benefit to exploring these paths.

# Example: Testing $I(\text{nedbarea}, \text{markgrm}, \text{dgv5980})$ in Barley

Any path through *aar\_mod* will eventually become **blocked**.



## DN Contribution

Stop traversing active paths that are doomed to become blocked.

BN	$ N $	d-Sep Tests	i-Sep Tests	Test Savings	Time Savings
Insurance	27	58898	36642	38%	-22%
Water	32	41392	23959	42%	-27%
Alarm	37	35224	23078	34%	-11%
Barley	48	78794	56804	28%	-15%
Hailfinder	56	51922	42543	18%	-23%
Pathfinder	135	125932	62820	50%	-79%
Munin1	186	167809	64329	62%	14%
Diabetes	413	827291	681468	18%	11%
Pigs	441	116795	12841	89%	36%
Link	724	336780	75505	78%	38%
Munin4	1038	509299	77314	85%	47%
Munin3	1041	459409	50147	89%	55%

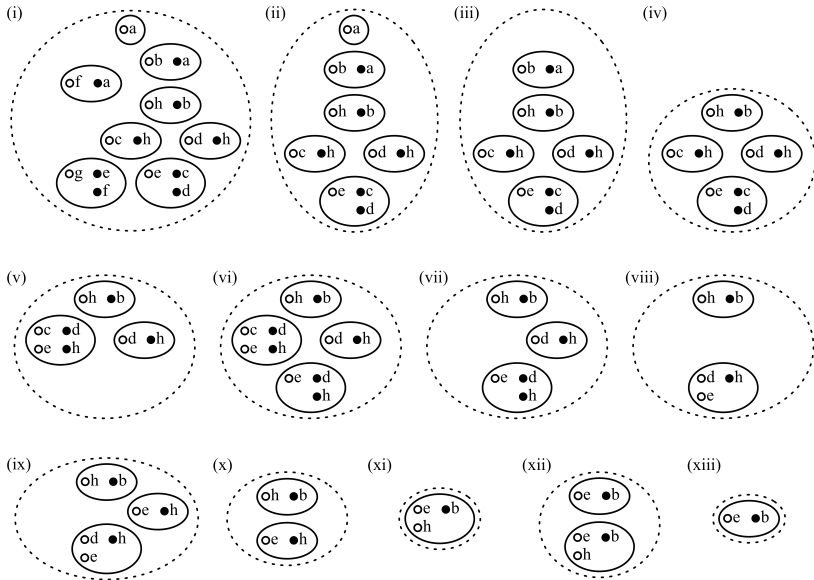
### 3. Advantages - Surprisingly Simple, Remarkably Robust

- In modeling, d-separation can use specialized terminology **not referenced in inference** such as “open sequential valves” and “closed divergent valves.”
- In inference, LP involves specialized terminology **not referenced in modeling** such as the “running intersection property.”
- DNs use the same terminology for inference and modeling.

## Representing All Steps of LP in DNs

- LP involves 2 networks.
- LP tests independencies in a BN, yet conducts inference in a JT.
- DNs do both in the same network.

# All Steps of VE: BN, query, independencies, computation



# Determining Good Elimination Orderings

- The order in which variables are eliminated can have a profound impact on the amount of computation performed.
- DNs can represent four well-known heuristics for determining good elimination orderings in BNs:
  - *min-neighbours* (MN),
  - *min-weight* (MW),
  - *min-fill* (MF),
  - *weighted-min-fill* (WMF).
- We introduced a new heuristic, called *potential energy* (PE), based on DNs.
- PE can score more accurately than the above heuristics.

## 4. Conclusion

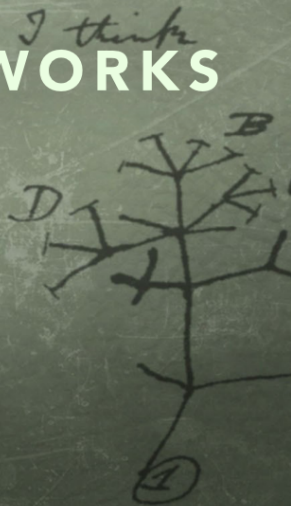
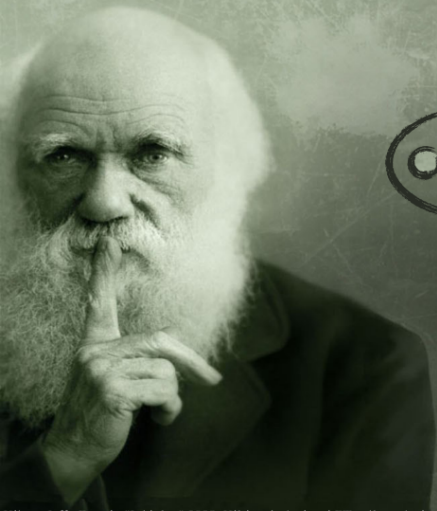
- **Purely graphical approach** to VE, which is often used to introduce BN inference to beginners
- **Faster way** to test independencies in BNs
- **Unify** modeling and inference into one network using common terminology
- DNs are like looking at BNs through a **microscope**

Watch [www.darwiniannetworks.com](http://www.darwiniannetworks.com) for updates



INTRODUCING

# DARWINIAN NETWORKS



*There between*