Bayesian Networks,

Mendelian Genetics,

and ...







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Blood Feud

In "Blood Feud" (S02E22):

- The rich Mr Burns badly needs
 O-blood for a transfusion
- Desesperate Mr Smithers asks nuclear power plant's employees to give blood for the great man





Homer Simpson sees a golden opportunity:

- Mr Burns being immensely rich, he will probably reward the one who saves him
- Homer asks Marge about his own blood group, but he's A+ (Homer: "D'oh!")

What are the chances to find a Oblood donor in the Simpson family?

The Blood Group/Rhesus system

Lisa asks Pr Frink about blood group genetics:

 ABO gene with three alleles: two codominant A, B, and one recessive O

$$\Rightarrow \textit{p}_{\textrm{O}} = 0.60, \, \textit{p}_{\textrm{A}} = 0.30, \, \textit{p}_{\textrm{B}} = 0.10$$

RHD gene with two alleles: dominant D (positive Rh) and recessive d (negative Rh)
 ⇒ q_D = 0.60, q_d = 0.40



With ABO/RHD independence we get:



| | ABO | 00 | OA | OB | AA | AB | BB |
|-----|--------------|------|------|------|------|------|------|
| RHD | ABO | 0.36 | 0.36 | 0.12 | 0.09 | 0.06 | 0.01 |
| DD | 0.36 | O+ | A+ | B+ | A+ | AB+ | B+ |
| Dd | 0.36 0.48 | O+ | A+ | B+ | A+ | AB+ | B+ |
| dd | 0.16 | O- | A- | B- | A- | AB- | B- |

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This leads to a total of 8 blood phenotypes:
 A+, B+, AB+, O+, A-, B-, AB-, O-





| | | group | 0 | Α | В | AB |
|---|----|-------|--------|-------|--------|--------|
| | Rh | | 0.36 | 0.45 | 0.13 | 0.06 |
| Ī | + | 0.84 | 0.3024 | 0.378 | 0.1092 | 0.0504 |
| | - | 0.16 | 0.0576 | 0.072 | 0.0208 | 0.0096 |



The problem to solve:

- 1: Homer, 2: Marge, 3: Bart, 4: Lisa, 5: Maggie
- X_i (resp. Y_i) genotype (resp. phenotype) or ind. i
- we need to compute

$$\pi = \mathbb{P}(\exists i, Y_i = \mathsf{O} - | Y_1 = \mathsf{A} +)$$

Model 1: independent genotypes:

$$\mathbb{P}(X, Y) = \prod_{i=1}^{5} \mathbb{P}(X_i) \mathbb{P}(Y_i | X_i)$$

- $\mathbb{P}(Y_1 = O | Y_1 = A +) = 0$, for $i \neq 1$, $\mathbb{P}(Y_i = O -) = 0.0576$
- $\pi = 1 (1 0.0576)^4 \simeq 0.2112$, easy, right!?

Lisa: "But genotypes are <u>not</u> independent!"

Homer: "D'oh!"



The problem to solve:

- 1: Homer, 2: Marge, 3: Bart, 4: Lisa, 5: Maggie
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- we need to compute

$$\pi = \mathbb{P}(\exists i, Y_i = \mathsf{O}\text{-}|Y_1 = \mathsf{A}\text{+})$$

Model 2: Mendelian transmission of alleles:

$$\mathbb{P}(X, Y) = \mathbb{P}(X_1)\mathbb{P}(X_2) \prod_{i=3}^{5} \mathbb{P}(X_i | X_1, X_2) \times \prod_{i=1}^{5} \mathbb{P}(Y_i | X_i)$$

•
$$\mathbb{P}(X_1 = \frac{\mathsf{OADd}}{\mathsf{OADd}} | Y_1 = \mathsf{A+}) = \frac{0.36}{0.36 + 0.09} \times \frac{0.48}{0.36 + 0.48} \simeq 0.4571$$

•
$$\mathbb{P}(X_2 = {}^{\mathsf{A}}_{\mathsf{B}}\mathsf{Dd}) = 0.48 \times 0.48 = 0.2304$$

•
$$\mathbb{P}(X_2 = O_B^A dd) = 0.48 \times 0.16 = 0.0768$$

•
$$\mathbb{P}(X_2 = \text{OODd}) = 0.36 \times 0.48 = 0.1728$$

•
$$\mathbb{P}(X_2 = \text{OOdd}) = 0.36 \times 0.16 = 0.0576$$



The problem to solve:

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Model 2: Mendelian transmission of alleles:

$$\mathbb{P}(X,Y) = \mathbb{P}(X_1)\mathbb{P}(X_2)\prod_{i=3}^{3}\mathbb{P}(X_i|X_1,X_2)\times\prod_{i=1}^{3}\mathbb{P}(Y_i|X_i)$$

 $ev = \{Y_1 = A+\}$ and N is the number of O- in the nuclear family

- $\mathbb{P}(X_1 = \mathsf{OADD}, \mathsf{AADd}, \mathsf{AADD}|\ \mathrm{ev}) = 0.5429 \Rightarrow \mathsf{N} \sim \mathcal{B}(1, 0.0567)$
- $\mathbb{P}(X_1 = \mathsf{OADd}, X_2 \text{ not carrier} | \mathsf{ev}) \simeq 0.2114 \Rightarrow \mathsf{N} = \mathsf{O}$
- $\mathbb{P}(X_1 = \mathsf{OADd}, X_2 = \mathsf{O}_\mathsf{B}^\mathsf{A}\mathsf{Dd}|\ \mathrm{ev}) \simeq 0.1053 \Rightarrow \mathsf{N} \sim \mathcal{B}(3, \frac{1}{16})$
- $\mathbb{P}(X_1 = \mathsf{OADd}, X_2 = \mathsf{O}_\mathsf{B}^\mathsf{A}\mathsf{dd}, \mathsf{OODd}|\ \mathrm{ev}) \simeq 0.1141 \Rightarrow N \sim \mathcal{B}(3, \frac{1}{8})$
- $\mathbb{P}(X_1 = \mathsf{OADd}, X_2 = \mathsf{OOdd}|\ \mathrm{ev}) \simeq 0.0263 \Rightarrow N \sim 1 + \mathcal{B}(3, \frac{1}{4})$



The problem to solve:

- 1: Homer, 2: Marge, 3: Bart, 4: Lisa, 5: Maggie
- X_i (resp. Y_i) genotype (resp. phenotype) or ind. i
- we need to compute

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Model 2: Mendelian transmission of alleles:

$$\mathbb{P}(X,Y) = \mathbb{P}(X_1)\mathbb{P}(X_2)\prod_{i=3}^{5}\mathbb{P}(X_i|X_1,X_2)\times\prod_{i=1}^{5}\mathbb{P}(Y_i|X_i)$$

 $ev = \{Y_1 = A+\}$ and N is the number of O- in the nuclear family

$$N | \text{ ev} \sim 0.5429 \times \mathcal{B}(1, 0.0567) + 0.2114 \times \mathbf{1}_{N=0} + 0.1053 \times \mathcal{B}(3, 1/16) + 0.1141 \times \mathcal{B}(3, 1/8) + 0.0263 \times (1 + \mathcal{B}(3, 1/4))$$

$$\sum_{n\geqslant 0}\mathbb{P}(\textit{N}=\textit{n}|\mathrm{ev})\textit{z}^{\textit{n}} \simeq 0.8867 + 0.0920\textit{z} + 0.0169\textit{z}^{2} + 0.0039\textit{z}^{3} + 0.0004\textit{z}^{4}$$

$$\pi = 0.0920 + 0.0169 + 0.0039 + 0.0004 = 0.1132$$

What does it mean?



Homer: "So $\pi = 0.1132$ or whatever, is that good?"

- $\mathbb{P}(\text{at least one O-}|\text{Homer is A+}) = 11.32\%$
- $\mathbb{P}(\text{not any O-}|\text{Homer is A+}) = 88.68\%$

Homer: "Huh ..."

9 to 10 chances of not being able to help Mr Burns

Homer: "D'oh!"

Frink: "But I found a blood test in the criminal

record of Bart ... "

Homer: "... and ?

Frink: "Bart is O-!!"

Homer: "Woohoo!"



Weak-D Bart

- Frink: "Alas, Bart has the weak-D phenotype. His blood might kill Mr Burns!"
- RHD gene, 3 alleles: D, d, and w (weak-D, pseudo neg Rh)

$$\Rightarrow q_{\rm D} = 0.60,\, q_{\rm d} = 0.39,\, q_{\rm w} = 0.01$$

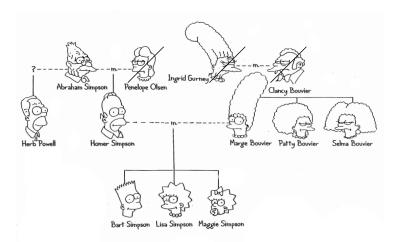
Only OOdd is compatible with Mr Burns!



| | ABO | 00 | OA | OB | AA | AB | BB |
|-----|--------|------|------|------|------|------|------|
| RHD | | 0.36 | 0.36 | 0.12 | 0.09 | 0.06 | 0.01 |
| DD | 0.3600 | 0+ | A+ | B+ | A+ | AB+ | B+ |
| Dd | 0.4680 | O+ | A+ | B+ | A+ | AB+ | B+ |
| Dw | 0.0120 | O+ | A+ | B+ | A+ | AB+ | B+ |
| dd | 0.1521 | O- | A- | B- | A- | AB- | B- |
| dw | 0.0078 | Ow | Aw | Bw | Aw | ABw | Bw |
| ww | 0.0001 | Ow | Aw | Bw | Aw | ABw | Bw |

Lisa: "We need to consider the extended Simpson family!"

Simpsons' Pedigree and Bayesian Network

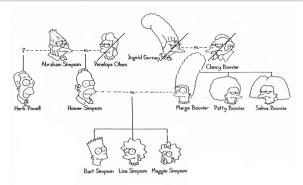


1: Herb's mother, 2: Abraham, 3: Penelope, 4: Ingrid, 5: Clancy,

6: Herb, 7: Homer, 8: Marge, 9: Patty, 10: Selma,

11: Bart, 12: Lisa, 13: Maggie

Simpsons' Pedigree and Bayesian Network



1: Herb's mother, 2: Abraham, 3: Penelope, 4: Ingrid, 5: Clancy, 6: Herb, 7: Homer, 8: Marge, 9: Patty, 10: Selma, 11: Bart, 12: Lisa, 13: Maggie

$$\begin{split} \mathbb{P}(X) &= \mathbb{P}(X_1) \mathbb{P}(X_2) \mathbb{P}(X_3) \mathbb{P}(X_4) \mathbb{P}(X_5) \\ \mathbb{P}(X_6 \mid X_{1,2}) \mathbb{P}(X_7 \mid X_{2,3}) \mathbb{P}(X_8 \mid X_{4,5}) \mathbb{P}(X_9 \mid X_{4,5}) \mathbb{P}(X_{10} \mid X_{4,5}) \\ \mathbb{P}(X_{11} \mid X_{7,8}) \mathbb{P}(X_{12} \mid X_{7,8}) \mathbb{P}(X_{13} \mid X_{7,8}) \end{split}$$

Simpsons' Pedigree and Bayesian Network

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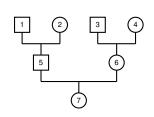
- $X = (X_1, X_2, \dots, X_{13})$ is the family genotype
- in order to compute $\mathbb{P}(\text{ev}) = \sum_{X'} \mathbb{P}(X', \text{ev})$
- we just have to sum over 81¹³ configurations

 $81^{13} = 6461081889226672446898176$

⇒ simply impossible!

Local computations in a simple pedigree

Idea: we consider a smaller (but similar) family, ev (evidence) still represents the available information.



• for founders (1, 2, 3, 4) i:

$$\varphi_i(X_i) = \mathbb{P}(X_i \cap \text{ev})$$

for offsprings (5, 6, 7)k with parents i, j:

$$\varphi_j(X_i, X_j, X_k) = \mathbb{P}(X_k \cap \text{ev} | X_i, X_j)$$



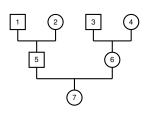
$$\mathbb{P}(\text{ev}) = \sum_{X_1} \sum_{X_2} \sum_{X_3} \sum_{X_4} \sum_{X_5} \sum_{X_6} \sum_{X_7} \varphi_1(X_1) \varphi_2(X_2) \varphi_3(X_3) \varphi_4(X_4)$$
$$\varphi_5(X_1, X_2, X_5) \varphi_6(X_3, X_4, X_6) \varphi_7(X_5, X_6, X_7)$$

 \Rightarrow 81⁷ = 22 876 792 454 961 still too large !!

Local computations in a simple pedigree

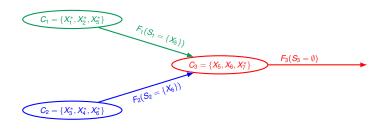
Pedigree

Clique decomposition



$$\mathbb{P}(\text{ev}) = \sum_{X_5} \sum_{X_6} \sum_{X_7} \left\{ \left(\sum_{X_1} \sum_{X_2} \varphi_1(X_1) \varphi_2(X_2) \varphi_5(X_1, X_2, X_5) \right) \right. \\ \left. \left(\sum_{X_2} \sum_{X_4} \varphi_3(X_3) \varphi_4(X_4) \varphi_6(X_3, X_4, X_6) \right) \varphi_7(X_5, X_6, X_7) \right\}$$

Local computations in a simple pedigree



$$F_{j}(S_{j}) = \sum_{C_{j} \setminus S_{j}} \left(\prod_{i \in \text{from}_{j}} F_{i}(S_{i}) \right) \times \prod_{X_{u} \in C_{j}^{*}} \varphi_{u}(X_{\text{pa}_{u}}, X_{u}) \qquad F_{3}(\emptyset) = \mathbb{P}(\text{ev})$$

Complexity:

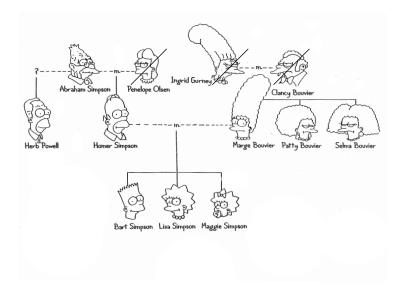
• from $81^7 = 22876792454961$

• to $3 \times 81^3 = 1594323$

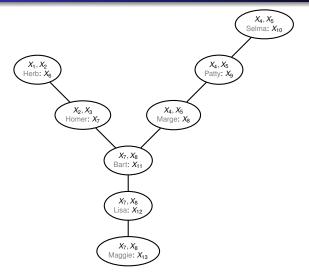
Lisa: "Much better!" Homer: "Woohoo!"



Clique decomposition for the Simpsons

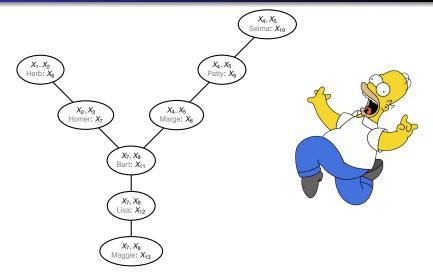


Clique decomposition for the Simpsons



- from $81^{13} = 6461081889226672446898176$
- to $8 \times 81^3 = 4251528$

Clique decomposition for the Simpsons



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Posterior Distribution

After the sum-product algorithm we can combine forward/backward quantities with the potentials φ_i in order to derive:

- the marginal distribution $\mathbb{P}(C_j|ev)$ of each clique;
- the marginal distribution $\mathbb{P}(S_j|ev)$ of each separator;
- the full posterior $\mathbb{P}(X|\text{ev})$ as a heterogenous Markov chain.

Introducing for all ind. *i* the *donor compatibility* random variable:

$$C_i = \mathbf{1}_{X_i = \mathsf{OOdd}} = \left\{ egin{array}{ll} 1 & ext{if } i ext{ compatible donor } i.e. \ X_i = \mathsf{OOdd} \\ 0 & ext{if } i ext{ not compatible donor } i.e. \ X_i
eq \mathsf{OOdd} \end{array}
ight.$$

Idea: introduce a dummy variable **z** in the sum-product s.t.

$$\sum_{n\geqslant 0}\mathbb{P}(N=n,\operatorname{ev})\mathbf{Z}^n=\sum_{X}\prod_{i}\varphi_i(X_{\operatorname{pa}_i},X_i)\mathbf{Z}^{\mathbf{1}_{X_i=\operatorname{OOdd}}}$$

$$\sum_{n\geqslant 0} \mathbb{P}(N=n|\text{ev})\mathbf{z}^n = 0.528 + 0.282\mathbf{z} + 0.0895\mathbf{z}^2 + 0.0404\mathbf{z}^3$$
$$+ 0.0329\mathbf{z}^4 + 0.0250\mathbf{z}^5 + 0.00189\mathbf{z}^6 + 0.000148\mathbf{z}^7 + 3.76 \times 10^{-7}\mathbf{z}^8$$

$$\mathbb{P}(C_i = 1 | \text{ev})$$

| | | • | | | |
|------------|---------|-------|--------|----------|--------|
| Herb's mum | Abrahaı | m Per | elope | Ingrid | Clancy |
| 5.5% | 2.4% | 2 | .4% | 16.2% | 16.2% |
| Herb | Homer | Marge | Patty | Selm | a |
| 3.8% | 0.0% | 13.5% | 10.9% | 10.99 | % |
| | Bart | Lisa | Maggie | <u> </u> | |
| | 0.0% | 1.5% | 1.5% | _ | |

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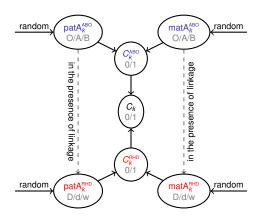
$$\sum_{n\geqslant 0}\mathbb{P}(\textit{N}=\textit{n},\text{ev})\textbf{Z}^\textit{n}=\sum_{\textit{X}}\prod_{\textit{i}}\varphi_{\textit{i}}(\textit{X}_{\text{pa}_{\textit{i}}},\textit{X}_{\textit{i}})\textbf{Z}^{\textbf{1}_{\textit{X}_{\textit{i}}}=\text{OOdd}}$$

$$\sum_{n\geqslant 0} \mathbb{P}(N=n|\text{ev})\mathbf{z}^n = 0.528 + 0.282\mathbf{z} + 0.0895\mathbf{z}^2 + 0.0404\mathbf{z}^3 + 0.0329\mathbf{z}^4 + 0.0250\mathbf{z}^5 + 0.00189\mathbf{z}^6 + 0.000148\mathbf{z}^7 + 3.76 \times 10^{-7}\mathbf{z}^8$$

| | | _ | Patty | |
|------|------|-------|--------|-------|
| 4.8% | 0.0% | 86.5% | 97.8% | 97.8% |
| | Bart | Lisa | Maggie | |
| | 0.0% | 3.2% | 3.2% | |

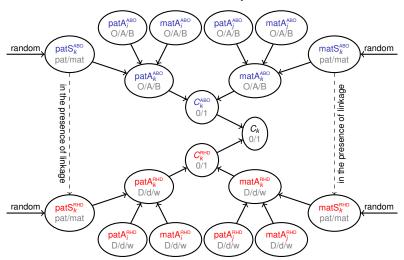
Extended Pedigree: Small Variables

For a *founder i*, instead of $X_i \in \mathcal{G}$ we have:



Extended Pedigree: Small Variables

For a *offspring* k (with father i and mother j), instead of $X_k \in \mathcal{G}|X_i, X_i$ we have:



Extended Pedigree: Small Variables

Recall on complexity:

- naive $81^{13} = 6461081889226672446898176$
- genotypes $8 \times 81^3 = 4251528$

Small variables with the three heuristics:

- min-neighbors: the smallest clique
 - ⇒ 61154 61649 89051
- min-fill: the clique with minimum fill-in
 - \Rightarrow 85205 92333 92360
- weighted min-fill: the clique with minimum weighted fill-in
 - ⇒ 57530 43841 43112



Epidemiological/Medical Applications

Many human diseases:

- Cancers:
 - Breast and Ovarian: Institut Curie
 - MSI Cancer and Lynch Syndrome: Saint-Antoine
 - Gliomas: La Pitié-Salpêtrière
- Rare Genetic Diseases:
 - Neuropathy Amyloid Hereditary: Henri Mondor
 - Pulmonary Arterial HT: Marie Lannelongue
 - Huntington Disease: Hôpital Saint-Anne
- Common Disease with Genetic Factors:
 - Alzheimer Disease: CHU Rouen
 - Diabetes, autism, cardio-vascular, obesity, . . .

And other applications: linkage analysis, genetic epidemiology, agronomy, recreative genetics, . . .



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- Matt Groening



