

DISTILLING CAUSAL MODELS MODEL AVERAGING, FEDERATED LEARNING AND MORE

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HERE COMES CLINICAL DATA

REALITY-CENTRIC CAUSAL NETWORKS

CONCLUSIONS

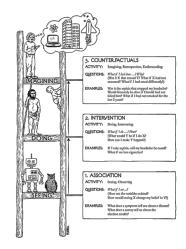
ACKNOWLEDGEMENTS

Machine learning creates black boxes that use probabilistic associations for prediction. Scientific questions are inherently causal.

We have [13, 21] a rigorous theory of causality that uses directed (acyclic) graphs (DAGs) to represent causes and effects. With it, we can reason about

- what we see,
- affecting change,
- hypothetical situations.

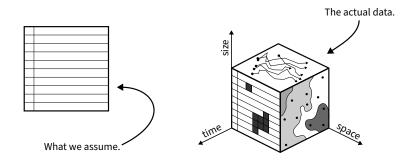
How can we learn causal DAGs from clinical data?



Learning a causal network means learning its structure \mathcal{G} and parameters Θ , much like Bayesian networks:



We used to ask domain experts for information [6, 7]; now we rely more and more on learning algorithms and the data \mathcal{D} [24].

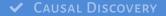


Bayesian networks [23] are defined by:

- a network structure, a directed acyclic graph (DAG) G, in which each node corresponds to a random variable X_i;
- a global probability distribution X with parameters Θ, which factorises into smaller local probability distributions according to the arcs in *G*:

$$\mathbf{P}(\mathbf{X}, \Theta) = \prod_{i=1}^N \mathbf{P}(X_i \mid \Pi_{X_i}; \Theta_{X_i}) \quad \text{where} \quad \Pi_{X_i} = \{\text{parents of } X_i\}.$$

Causal networks are Bayesian networks with extra assumptions, such as causal sufficiency, that allow for interpreting arcs as causal effects.



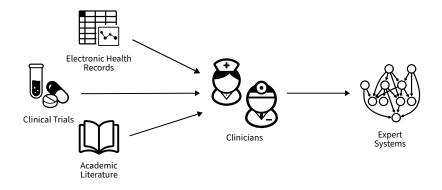
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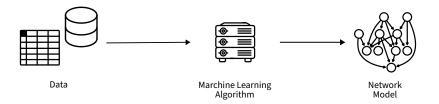
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First, there were expert systems [3, 4] compiled with clinicians.



The clinicians distilled all the available knowledge into a set of causal effects (\mathcal{G}) and their effect sizes (Θ).

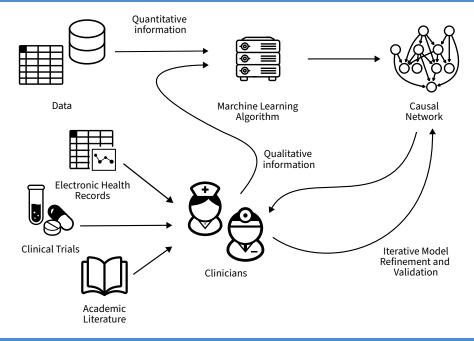
However, clinicians cannot scale to complex phenomena and often do not agree with each other. Machine learning then took a completely data-driven approach.



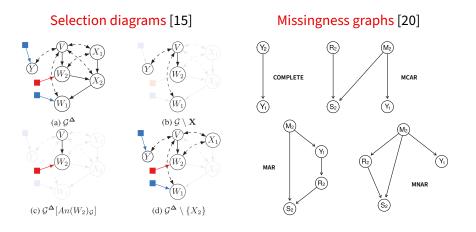
This is the current paradigm of structure learning (for Bayesian networks) and causal discovery (for causal networks) [14, 28]. Many, many such algorithms in the literature [24].

- We do not have enough data to capture the complexity of modern clinical problems.
- Causal discovery algorithms have unrealistic assumptions and are (mostly) unusable on clinical data.
- Most of the field has forgotten the accumulated statistical wisdom on the design of experiments, hierarchical data, model validation, etc.
- There is an excess focus on artificial benchmarks that have nothing to do with the challenges and the risk evaluations of real clinical practice.
- The expertise of clinicians is under-appreciated and under-utilised in driving causal discovery.

PRODUCING REALISTIC, USEFUL MODELS

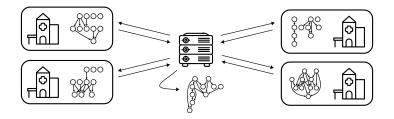


Causal networks require causal sufficiency: no latent confounders. Both sampling bias and incomplete data are such if left unmodelled.



How do we aggregate data across centres when we cannot share it?

- Score-based: computing local scores for parent sets in the centres and aggregating them to minimise regret [18, 19].
- Constraint-based: aggregating partial networks learned using conditional independence tests [12].



The "true network" may be different in each centre, but causal discovery assumes that there is a single one that is valid for all centres...

Individuals are not independent and identically distributed!

- Panel data are correlated over time.
- Epidemiological data are correlated over space.
- Subgroups of patients with differential treatment effects.
- Centre heterogeneity is unavoidable in multi-centre clinical trials.

And yet mostly $X_i = f(\Pi_{X_i}; \Theta_{X_i}) + \varepsilon_i$ with $\operatorname{COV}(\varepsilon_{i,j}, \varepsilon_{i,k}) = 0$.

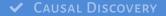
- We need hierarchical models sharing information between centres, as in [1, 27], or to borrow ideas from federated learning.
- We definitely need to borrow models for state-space data from the ecological modelling literature.

MLOps [26] makes an important distinction:

- Model evaluation: is the statistical performance good?
- Model validation: what about the domain metrics we care about?

Machine learning only cares about the former, and even then there is no consensus on how to do that.

- Many existing metrics fail to exhibit a strong correlation with the quality of the approximation to the true model posterior [17].
- Different causal structural metrics with contrasting interpretations
 [11]. SID [22] is inconsistent and very costly to compute.
- Even quantitative metrics such as [5] are not framed in clinical terms (say, ATE). An exception may be [8].



✓ HERE COMES CLINICAL DATA

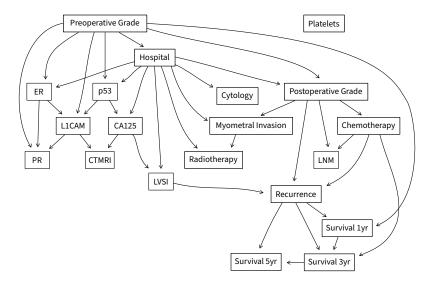
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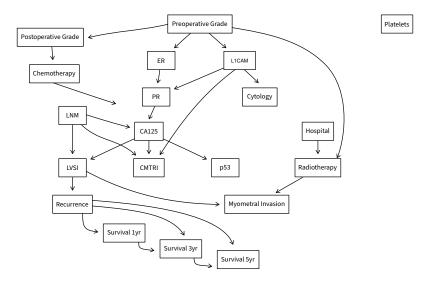
ACKNOWLEDGEMENTS

- Ten clinics in the European Network for Individualised Treatment of Endometrial Cancer (ENITEC).
- Key Issues:
 - Small sample size.
 - High missingness ratio.
 - Centre heterogeneity.
- Modelling:
 - Prior knowledge from clinicians.
 - Model averaging.
 - MNAR missingness.

Radiostreenand und cruster radiosed	p53	Recurrence	Primary tumor	Platelets	PR	Myometral invasion	LNM	LICAM	ER	Cytology	CTMRI	CA125	Chemotherapy	Radiotherapy
ienoth				0.1									1.0	
Etapy .				0.1										
ALLS	0.3		0.3	0.3				0.1		0.4	0.2			
CIMA	-0.1			0.4	0.1		0.2	-0.1						
JOEN	0.3		0.4	0.1	0.1		-0.1	0.1						
~			0.1											
L'CAM	-0.1		0.1				-0.1							
MM	-0.1		-0.2	0.3										
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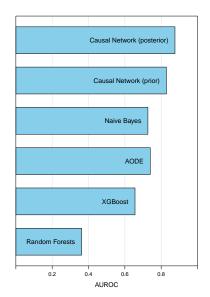


Baseline model learned with SEM [9, 10] from our earlier AlxIA paper [29].

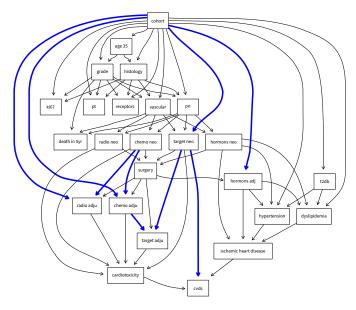


Unbiased model learned with HC-aIPW [16] in our AIME paper [30].

- A population cohort and a clinical cohort from Istituto Nazionale dei Tumori di Milano.
- Key Issues:
 - Different sampling criteria.
 - Different sets of variables.
 - Low prevalence / incidence.
- Modelling:
 - Prior knowledge from clinicians.
 - Model averaging.
 - MNAR missingness.
 - Sampling diagrams.



MODELLING MULTIPLE COHORTS TOGETHER



Model learned with SEM in our AIxIA paper [2].

Google

Google COVID-19 Open Data: 400 health conditions, 4 countries (county-level in the US), weekly search frequencies for 2020-2023 normalised by NLP. Weather stations in 1652 counties with and satellite images.





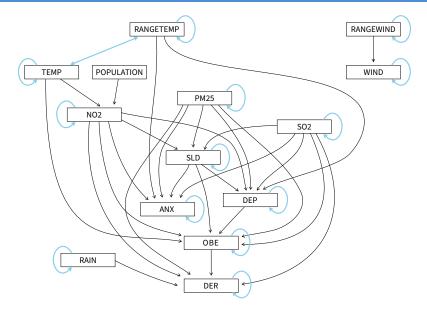
Monitoring stations in 1470 counties with hourly measurements of NOx, SOx, O3, PMx.

Socio-economic data at the population level to avoid confounding.



Key Issues: incomplete and heterogeneous state space-data (\approx 53k observations over \approx 500 US counties and 134 weeks.) Modelling: GLS, IRLS, node-averaged likelihood, model averaging.

DISREGARDING THE STATE-SPACE NATURE OF THE DATA



Biased model learned with a dynamic network as in my previous work [25].

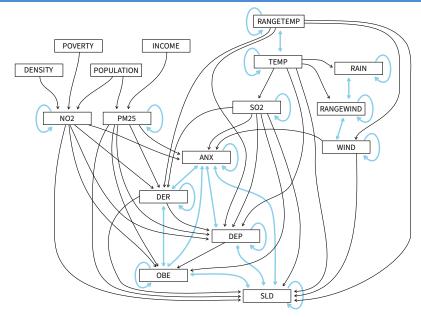
Residuals are reasonably free from autocorrelation, but they are full of spatial correlation and heteroscedastic!

	lag 1	lag 2	lag 3	lag 4	space	heteroscedastic
ANX	0.024	0.000	0.000	0.048	0.460	$4 imes 10^{ ext{-169}}$
DEP	0.016	0.000	0.000	0.000	0.325	$1 imes 10^{-212}$
DER	0.032	0.000	0.000	0.000	0.754	0
OBE	0.000	0.000	0.000	0.000	0.563	$6 imes10^{ ext{-100}}$
SLD	0.092	0.007	0.007	0.000	0.381	$1 imes 10^{-154}$

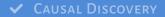
Reality chooses a better model:

- Adding a spatial correlation matrix: BF = 81.59.
- Adding different standard errors for different states: BF = 25.31.
- Reducing sparsity to let known arcs in: BF = 72.15.

MODELLING THE STATE-SPACE NATURE OF THE DATA



Unbiased model that I am currently working to improve.



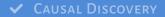
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- Causal networks are a principled and versatile tool to model clinical data in their complexity.
- Statistical and clinical experts are fundamental: using causal networks as a purely machine learning model is doomed to fail.
- State-space data, mixed variable types, missing values, population structure, non-stationarity: we can deal with them!
- There are technical challenges in translating clinical requirements and knowledge into causal discovery approaches, but we are actively working on them.



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→ ACKNOWLEDGEMENTS

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THAT'S ALL!

HAPPY TO DISCUSS IN MORE DETAIL.



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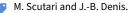
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