## Inference of Cause and Effect with Unsupervised Inverse Regression

## Abstract

We address the problem of causal discovery in the two-variable case, given a sample from their joint distribution. Since  $X \to Y$  and  $Y \to X$  are Markov equivalent, conditional-independence-based methods [Spirtes et al., 2000, Pearl, 2009] can not recover the causal graph. Alternative methods, introduce asymmetries between cause and effect by restricting the function class (e.g., [Hoyer et al., 2009]).

The proposed causal discovery method, CURE, is based on the principle of independence of causal mechanisms [Janzing and Schölkopf, 2010]. For the case of only two variables, it states that the marginal distribution of the cause, say P(X), and the conditional of the effect given the cause P(Y|X) are "independent", in the sense that they do not contain information about each other (informally P(X) " $\bot$ " P(Y|X)). This independence can be violated in the backward direction: the distribution of the effect P(Y) and the conditional P(X|Y) may contain information about each other because each of them inherits properties from both P(X) and P(Y|X), hence introducing an asymmetry between cause and effect. For deterministic causal relations (Y = f(X)), all the information about the conditional P(Y|X) is contained in the function f, so independence boils down to P(X) " $\bot$ " f. Previous work formalizes the independence principle by specifying what is meant by independence. For deterministic non-linear relations, Janzing et al. [2012] and Daniusis et al. [2010] define independence as uncorrelatedness between  $\log f'$  and the density of P(X), both viewed as random variables. For non-deterministic relations, it is not obvious how to explicitly formalize independence between P(X) and P(Y|X). Instead, we propose an implicit notion of independence, namely that  $p_{Y|X}$  cannot be estimated based on  $p_X$  (lower case denotes density). However, it may be possible to estimate  $p_{X|Y}$  based on the density of the effect,  $p_Y$ .

In practice, we are given empirical data  $\mathbf{x} \in \mathbb{R}^N$ ,  $\mathbf{y} \in \mathbb{R}^N$  from P(X, Y) and estimate  $p_{X|Y}$  based on  $\mathbf{y}$  (intentionally hiding  $\mathbf{x}$ ). The relationship between the observed  $\mathbf{y}$  and the *latent*  $\mathbf{x}_{\mathbf{u}} \in \mathbb{R}^N$  is modeled by a Gaussian Process (GP):  $p(\mathbf{y}|\mathbf{x}_{\mathbf{u}}, \boldsymbol{\theta}) = \mathcal{N}(\mathbf{y}; \mathbf{0}, K_{\mathbf{x}_{\mathbf{u}}, \mathbf{x}_{\mathbf{u}}} + \sigma_n^2 I_N)$  (this can be alternatively seen as a single output GP-LVM). Then, the required conditional  $p_{X|Y}$  is estimated as  $\hat{p}_{X_u|Y}^{\mathbf{y}}$ :  $(x_u, y) \mapsto p(x_u|y, \mathbf{y})$ , with  $p(x_u|y, \mathbf{y})$  estimated by marginalizing out the latent  $\mathbf{x}_{\mathbf{u}}$  and  $\boldsymbol{\theta}$  (GP hyperparameters).

CURE infers the causal direction by using the procedure above two times: one to estimate  $p_{X|Y}$  based only on y and another to estimate  $p_{Y|X}$  based only on x. If the first estimation is better,  $X \to Y$  is inferred. Otherwise,  $Y \to X$ . CURE was evaluated on synthetic and real data and often outperformed existing methods. On the downside, its computational cost is comparably high. This work was recently published at AISTATS 2015 [Sgouritsa et al., 2015].

## References

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