Controlling Selection Bias in Causal Inference

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Abstract
Selection bias, caused by preferential exclusion of units (or samples) from the data, is a major obstacle to valid causal inferences, for it cannot be removed or even detected by randomized experiments. This paper highlights several graphical and algebraic methods capable of mitigating and sometimes eliminating this bias. These nonparametric methods generalize and improve previously reported results, and identify the type of knowledge that need to be available for reasoning in the presence of selection bias.

Introduction
Selection bias is induced by preferential selection of units for data analysis, and is often governed by unknown factors including treatment, outcome and their consequences. Case-control studies in Epidemiology are particularly susceptible to such bias, e.g., cases may be reported only when the outcome (disease or complication) is unusual, while non-cases remain unreported (see (Glymour and Greenland 2008; Robins, Herman, and Brumback 2000; Hernán, Hernández-Díaz, and Robins 2004))

To illustrate the nature of this bias, consider the model of Fig. 1 (a) in which $S$ is a variable affected by both $X$ and $Y$, indicating entry into the data pool. Such preferential selection to the pool amounts to conditioning on $S$, which creates spurious associations between $X$ and $Y$ through two mechanisms. First, conditioning on $S$ induces spurious association between its parents, $X$ and $Y$. Second, $S$ is also a descendant of a “virtual collider” $Y$, whose parents are $X$ and the error term $U_Y$ (representing “omitted factors”) which is always present, though often not shown in the diagram.

A medical example of selection bias was reported in (Horwitz and Feinstein 1978), and subsequently studied in (Hernán, Hernández-Díaz, and Robins 2004; Geneletti, Richardson, and Best 2009), in which it was noticed that the effect of Oestrogen ($X$) on Endometrial Cancer ($Y$) was overestimated in the data studied. One of the symptoms of the use of Oestrogen is vaginal bleeding ($W$) (Fig. 1(c)) and the hypothesis was that women noticing bleeding are more likely to visit their doctors, causing women using Oestrogen to be overrepresented in the study.

Methods for controlling selection bias are relevant because such bias is pervasive in almost all empirical studies, including Machine Learning, Bioinformatics, Epidemiology and Economics.

Contributions
Our contributions are described in full details in the extended version of this paper (Bareinboim and Pearl 2011); here we highlight our main results.

Assuming no confounding, we quantify causal effects by the odds ratio $OR(Y, X \mid Z = z) = (Pr(y \mid z, x')/Pr(y' \mid z, x'))/(Pr(y \mid z, x)/Pr(y' \mid z, x))$. One of the properties of the odds ratio is that it is symmetric, i.e., $OR(X, Y \mid Z) = OR(Y, X \mid Z)$; another property is that it does not depend on the marginal distributions. It is well known (Cornfield 1951; Geng 1992) that the odds ratio is recoverable, i.e., $OR(X, Y \mid Z, S = 1) = OR(X, Y \mid Z)$ if either ($X \perp S \mid \{Y, Z\}$) or ($Y \perp S \mid \{X, Z\}$). One such example is depicted in Fig. 2(a), with $Z = \emptyset$.

In Fig. 2(b) the $OR(X, Y \mid Z)$ is recoverable but devoid of any causal interpretation. This is so because it does not stand for a causal effect in a a stable subset of individuals. Since $Z$ is $X$-dependent in $G$, the class of units for which $Z = z$ under $do(X = 1)$ is not the same as the class of units for which $Z = z$ under $do(X = 0)$. The conditional odd ratio $OR(X, Y \mid Z)$ would be meaningful only if $Z$ is restricted to pre-treatment covariates, which are $X$-invariant, hence stable.

By contrast, in the graph in Fig. 2(c) the $OR(X, Y \mid C)$ is recoverable and meaningful, and equates to

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Figure 1: (a,b) Simplest examples of selection and confounding bias, respectively. (c) Typical study with intermediary variable $W$ between $X$ and selection.

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1See (Pearl 2009, pp. 339-341) for further explanation of this bias mechanism.
Theorem 1 (Bareinboim and Pearl 2011, Thm. 2) Let graph $G$ contain the arrow $X \rightarrow Y$ and a selection node $S$. A necessary and sufficient condition for $G$ to permit the $G$-recoverability of $OR(Y, X \mid C)$ for some set $C$ of pre-treatment covariates is that every ancestor $A_i$ of $S$ that is also a descendant of $X$ have a separating set $T_i$ that either $d$-separates $A_i$ from $X$ given $Y$, or $d$-separates $A_i$ from $Y$ given $X$. (We define $S$ to be among its ancestors).

Moreover, if the condition above holds, then $OR(Y, X \mid C)$ is $G$-recoverable if $C$ $d$-separates $Y$ (given $X$) from all pre-treatment variables that are invoked in at least one $T_i$, and it equals $OR(Y, X \mid C, T, S = 1)$, where $T = \bigcup_i T_i$.

For instance, the problem studied in (Geneletti, Richardson, and Best 2009) (Fig. 1(c)) is trivially solved by the Theorem 1, yielding $OR(X, Y) = OR(X, Y \mid W, S = 1)$.

Having characterized recoverability of the odds ratio, we studied universal curves that show the behavior of OR as the distribution $P(y \mid x)$ changes, and how other measures of association such as risk ratio (RR) and risk difference (RD) are related to OR. We further showed that if one is interested in recovering RR and RD under selection bias, knowledge of the marginal distribution $P(X)$ is sufficient for recovery.

The results stated so far focus on point identifiability for the quantity of interest, but there are abundant scenarios where causal effects are non-identifiable and selection bias can simultaneously be present, further increasing the bias. One of the methods used to cope with non-identifiability is to bound the causal effects through instrumental variables (Pearl 2009, Ch. 8), but since the bounding analysis assumes no selection bias, the question arises whether tighter bounds can be derived in the presence of selection bias.

We show that selection bias can be removed entirely through the use of IVs, therefore, the bounds on the causal effect will be narrower than those obtained under the selection-free assumption as follows:

Theorem 2 (Bareinboim and Pearl 2011, Corol. 3) The bounds for the causal effect of $X$ on $Y$ can be recovered from selection bias whenever the following conditions hold:

(i) the $S$ node is affected by the set $Z$ only through $\{X, Y\}$;

(ii) the set $Z$ is $d$-connected to $\{X, Y\}$ (and combinations);

(iii) the dimensionality of $Z$ matches the dimensionality of $\{X, Y\}$; (iv) the marginal distribution of $Z$ is known.

This result is surprising for two reasons: first, we generally do not expect selection bias to be removable; second, bias removal in the presence of confounding is generally expected to be a more challenging task. We finally show how this result is applicable to scenarios where other structural assumptions hold, for instance, when an instrument is not available but a certain back-door admissible set can be identified (Bareinboim and Pearl 2011, Corol. 3–5).

Conclusion

We showed that qualitative knowledge of the selection mechanism together with graphical and algebraic methods can eliminate selection bias in many realistic problems. In particular, the paper provides a simple graphical condition, together with an algorithm to decide, given a DAG with measured and unmeasured variables, whether and how a given $e$-specific odds ratio can be recovered from selection-biased data. We further showed by algebraic methods that selection bias can be removed in the presence of confounding with the help of instrumental variables under certain mild conditions.

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References


