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Automated selection of Negative Control Exposure-Outcome pairs for use in Observational Studies: A capabilities demonstration

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Abstract

Negative control exposure-outcome pairs should be more commonly employed in observational studies to reduce the potential for systematic error^{1,2}. When employed, negative controls help to identify and calibrate for confounding as well as other sources of error. The process for identifying negative controls (either exposure controls or outcome controls) for use in observational studies requires time-consuming, manual curation of information from various knowledge sources. Here we demonstrate a means for automating the selection of candidate negative controls by leveraging an open-source standardized evidence base known as LAERTES (Large-scale Adverse Effects Related to Treatment Evidence Standardization) made available through the Observational Health Data Sciences and Informatics (OHDSI) collaborative.

Introduction

The LAERTES evidence base integrates several sources of evidence for investigating the association between drugs and health outcomes of interests (HOIs) into a single data source. Evidence is sourced from spontaneous reports, scientific literature, and both American and European product labeling. A statistical model was created using the evidence in LAERTES for discriminating between known positive drug-HOI causal relationships and drugs known to be unassociated with an outcome.

ATLAS is an open-source platform made available through OHDSI that provides a single user-experience for generating evidence. ATLAS has been integrated with LAERTES to provide a workflow for automatically identifying positive and negative controls.

Methods

The LAERTES evidence base contains information from spontaneous reports, scientific literature, and product labeling. Spontaneous reporting evidence is culled from the FDA Adverse Event Reporting System (FAERS) and includes counts of reports and proportional reporting ratio (PRR) scores^{3,4}. Evidence from the scientific literature was generated through two methods: one leveraging Medical Subject Headings (MeSH) tags following the process described by Avillach et al.⁵, and another (SemMedDB) that uses relationships semantically tagged within Medline abstracts natural language processing⁶. The results of these two methods are additionally stratified by Medline publication types: clinical trials, case report, and all other abstracts (i.e., of type Meta-Analysis, Comparative Study, Multicenter Study, or Journal Article). Finally, American product labels are parsed by a method developed by Duke et al.⁸, and ADRs mentioned in European labels are provided by the PROTECT project⁷.

The evidence in LAERTES was narrowed down to a “universe” of drugs and HOIs that met a specific, minimum criterion of evidence. The minimum criteria is defined as: the condition or ingredient had at least one FAERS evidence item, one Medline evidence item and one evidence item from a product label. This step was taken because a condition or drug that did not meet the minimum criteria might indicate a lack of clinical experience.

Logistic regression was used to build a multivariable prediction model on the LAERTES data that could discriminate between positive and negative controls. Regularization with a Laplace prior on the regression coefficients was used to allow the model to perform parameter selection. A union of the OMOP Reference Set⁹ and the Exploring and Understanding Adverse Drug Reactions (EU-ADR) Reference Set¹⁰⁻¹² was used to train our classifier. The prediction accuracy was estimated using leave-pair-out cross-validation.

An automated process was developed to expose the prediction model along with the evidence “universe” to return positive and negative controls. The input to the algorithm is a list of OMOP concept ids that define either a drug ingredient or an HOI. When given a drug ingredient, the algorithm will return the full “universe” of HOIs with counts for each evidence source and the predicative values. It works in a similar fashion when given an HOI, the algorithm will return the same information for all drug ingredients in the “universe”.

Results

Figure 1 provides an example of input to the negative controls algorithm while Figure 2 shows the output including the evidence counts, by source and the prediction values.

Concept Id	Concept Code	Concept Name	Domain	Standard Concept Caption	Exclude	Descendants	Mapped
192671	74474003	Gastrointestinal hemorrhage	Condition	Standard	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>

Figure 1. Input to the negative control algorithm (HOI: Gastrointestinal hemorrhage)

Id	Name	Domain	Medline CT	Medline Case	Medline Other	Splicer ADR	AERS	AERS PRR	Prediction	RC	DRC
714785	Amphetamine	DRUG	0	0	0	0	24	1.2208	0.0540	1,626,031	8,512,293
19069022	Sodium Fluoride	DRUG	0	0	0	0	18	0.8166	0.0540	929,189	2,291,729
1130863	Brompheniramine	DRUG	0	0	1	0	18	0.9203	0.0540	740,564	1,570,712
1101703	homatropine	DRUG	0	0	0	0	4	2.8242	0.0540	714,708	1,549,585
1519936	Etonogestrel	DRUG	0	0	0	0	33	0.2274	0.0540	587,107	2,802,049
967562	lorteprednol etabonate	DRUG	0	0	0	0	11	0.3452	0.0540	382,574	827,289
1036525	Sulfur	DRUG	0	0	0	0	19	3.7349	0.0540	357,668	816,784
1189697	eletriptan	DRUG	0	0	1	0	23	0.4735	0.0540	354,471	1,321,070
731533	dexamethylphenidate	DRUG	0	0	0	0	2	0.1724	0.0540	276,147	1,317,112
732893	Bupivacaine	DRUG	0	0	0	0	26	0.3053	0.0540	267,887	267,960
922868	Permethrin	DRUG	0	0	2	0	5	0.7223	0.0540	241,221	525,136
1158632	Methscopolamine	DRUG	0	0	0	0	6	5.2432	0.0540	197,867	456,217
1551673	Estrogens, Esterified (USP)	DRUG	0	0	1	0	22	0.5004	0.0540	194,968	1,045,003
1505346	Triiodothyronine	DRUG	0	0	0	0	17	0.1783	0.0540	163,902	819,723
951279	Prilocaine	DRUG	0	0	0	0	9	1.6126	0.0540	159,242	349,570

Figure 2. Output of the negative control algorithm: drug ingredients in the “universe”, their evidence counts and predictive value.

Conclusion

Automating a list of candidate negative controls using LAERTES and ATLAS lowers the amount of manual effort required of researchers while implementing a study.

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