

# ZaliQL: Causal Inference from Observational Data at Scale

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## ABSTRACT

Causal inference from observational data is a subject of active research and development in statistics and computer science. Many statistical software packages have been developed for this purpose. However, these toolkits do not scale to large datasets. We propose and demonstrate ZaliQL: a SQL-based framework for drawing causal inference from observational data. ZaliQL supports the state-of-the-art methods for causal inference and runs at scale within PostgreSQL database system. In addition, we built a visual interface to wrap around ZaliQL. In our demonstration, we will use this GUI to show a live investigation of the causal effect of different weather conditions on flight delays.

## 1. INTRODUCTION

Randomized experiments (AB testing) remain the gold standard for causal inference; however, they do pose a number of problems. Namely, controlled experiments are not feasible for ethical, economical, or practical reasons in a number of disciplines [9]. Observational studies can be used to draw causal inference without controlled experiments [9, 11].

Computational, physical, and social scientists all increasingly want to perform causal inference on big datasets from observational studies including social networks, biological networks, sensor networks, and others. Unfortunately, the current software for processing observational data in terms of causal inference simply cannot scale. R, Stata, SAS, and SPSS all have packages; however, they are designed to be used only with single table data, making them cumbersome and often ineffective with large datasets [4, 6]. For example, we found performing CEM on a dataset with 5M entries takes up to an hour using Stata, R, or SAS. This is obviously not an effective practice for researchers.

Additionally, causal analysis is part of a larger pipeline that includes data acquisition, cleaning, and integration. For large datasets, these tasks are better handled by a relational database engine which provides most of the functionality needed for these tasks while also scaling up to large datasets.

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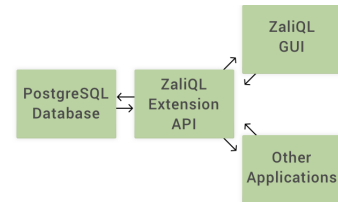


Figure 1: ZaliQL architecture

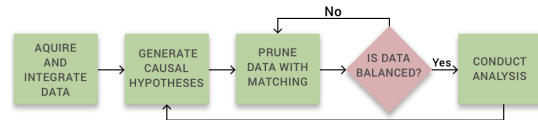


Figure 2: Causal analysis workflow

The purpose of this demonstration is to propose ZaliQL1,<sup>1</sup> an SQL-based framework for drawing causal inference within the DBMS. ZaliQL takes the initial step towards truly scalable causal inference by modeling it as a data management problem. We demonstrate that causal inference can be approached from this perspective and that doing so is key to scalability and robustness of the data. ZaliQL supports innovative methods for causal inference and runs at scale within a database engine.

## 2. SYSTEM ARCHITECTURE

The overall architecture of ZaliQL can be seen in Fig. 1. The API is a set of functions that support causal inference on data stored in a PostgreSQL DBMS. The API will be packaged as a PostgreSQL extension. The ZaliQL API is modeled after the MatchIt and CEM toolkits [4, 6] and includes methods for drawing causal inference from relational data. A web GUI is also included for demonstration and exploration purposes and is illustrated in Fig. 3.

## 3. DEMONSTRATION DETAILS

As the illustration in Fig. 2 shows, modern causal analysis is an iterative process. An analyst must acquire and integrate data from a myriad of sources, generate a hypothesis, pre-process the data through a matching method, and finally conduct the causal analysis. This linear process often must be repeated with new matching methods, new hypotheses,

<sup>1</sup> The prefix Zali refers to al-Ghazali (1058-1111), a medieval Persian philosopher. It is known that David Hume (1711-1776), a Scottish philosopher, who gave the first explicit definition of causation in terms of counterfactuals, was heavily influenced by al-Ghazali's conception of causality.

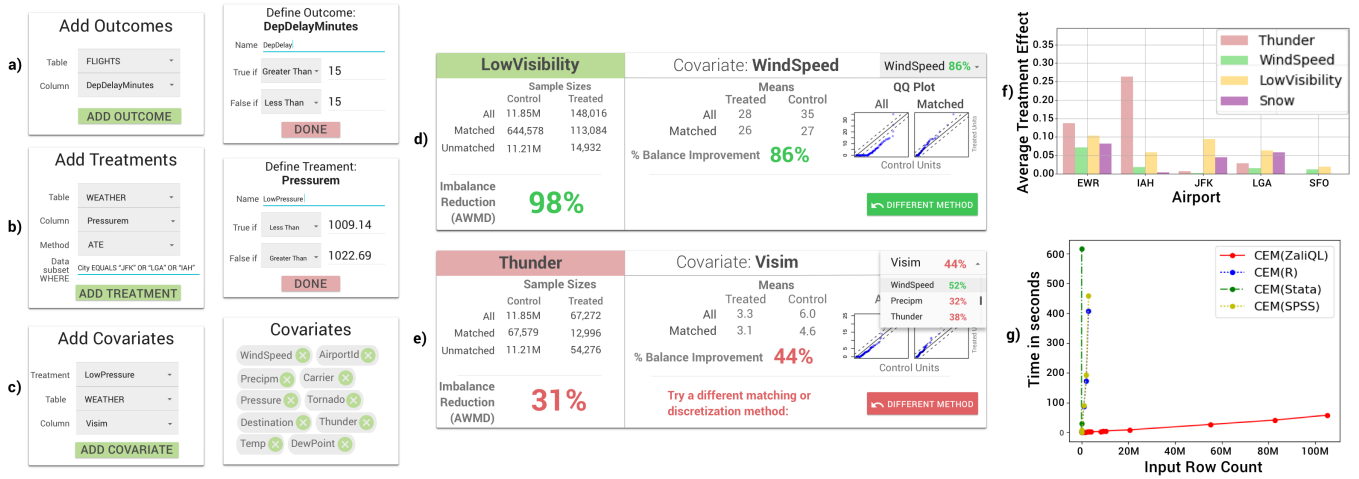


Figure 3: Demonstration screenshot described in Section 3

or even new datasets [5]. We will demonstrate ZaliQL by providing walkthroughs using causal investigations on integrated flight and weather datasets.

**Data:** The analysis will be conducted on a spatio-temporal join of the following datasets: (a) Flight dataset (105M entries) collected by the U.S. Department of Transportation.<sup>2</sup> This dataset contains records of over 90% of U.S. domestic flights of major airlines between 1988 and the present. It includes the following variables: FlightDate, OriginAirportID, CarrierID, CRSDepTime (scheduled departure time), and DepDelay (departure delay); (b) *Weather dataset (40M entries)*: collected using Weather Underground API.<sup>3</sup> It contains historical weather data that is related to flights and includes the following variables: Code (Airport ID), Date, Time, Visim (visibility in km), Tempm (Temperature in Celsius), Wspdmm (wind speed in kph), Pressurem (Pressure in mBar), Precipm (Precipitation in mm), Snow (binary), and Thunder (binary).

**Data exploration:** Our demonstration will start by exploring the effect of different weather features on flight departure delay. As an example, we will show that 11% of flights were delayed when pressure was low; however, only 0.4% of flights were delayed when pressure was high. This suggests that pressure is inversely correlated with flight delay. However, after grouping by different variables such as airport, airline, and other weather features, flight delay frequency declined from 11% to nearly 0%.<sup>4</sup> Thus, we are forced to reconsider if low pressure is really a causative factor for weather delays.

**Causal questions:** The following causal questions have been defined: Q1: Does low air pressure cause flight departure delays? Q2: Which weather features are major causes of departure delays? Q3: Do the findings to the previous questions differ between major airports? These questions will be answered by ZaliQL using the following steps:

- Specifying DepDelay as our outcome of interest (effect) as shown in Fig. 3a.
- Specifying a set of binary treatments (causes) that could affect DepDelay as shown in Fig. 3b. Particularly, the following binary treatments will be created: LowVisibility (1 if Visim < 1); HeavySnow (1 if Precipm > 0.3 and Snow = 1); HighWindSpeed (1 if Wspdmm > 40; 0 if Wspdmm < 20); Thunder; Low-Pressure (1 if Pressurem < 1009.14; 0 if Pressurem > 1022.69).
- Specifying a subset of data that is relevant to the analysis. We will select five U.S. airports with high rates of weather-related delay. Specifically, San Francisco (SFO), John F. Kennedy (JFK), Newark Liberty (EWR), George Bush-Houston (IAH), and LaGuardia (LGA).

**Computing ATE:** The primary objective of causal inference is to quantify the causal effect of a binary treatment on an outcome. As a result, a common measure of effect is average treatment effect (ATE). ATE is defined as the difference in expected values of the treated and untreated units.<sup>5</sup> For example, regarding Q1, ZaliQL computes ATE as:

$$\mathbb{E}[\text{DepDelay} | \text{LowPressure} = 1] - \mathbb{E}[\text{DepDelay} | \text{LowPressure} = 0]$$

In this example,  $\mathbb{E}[\text{DepDelay} | \text{LowPressure} = x]$ ,  $x = (0, 1)$  is computed by taking the emetical average of DepDelay where LowPressure is  $x$ . It is quite common to compare an outcome (DepDelay) in the *treated group*, those subjects (flights) that receive a treatment (LowPressure = 1) with the *control group* (LowPressure = 0). We show that for LowPressure, ATE is 10 minutes. This is relatively large and suggests that pressure affects DepDelay. However, it is known from prior analysis that LowPressure alone does not serve as the sole causative factor for departure delay. This observation thus raises the question: Where is this difference coming from?

**Confounding variables:** The difference is a product of confounding variables, which make it difficult to establish a

<sup>2</sup><http://www.transtats.bts.gov/>

<sup>3</sup><https://www.wunderground.com>

<sup>4</sup>This phenomenon is known as Simpson's paradox and arises frequently in observational studies [8]. We demonstrate that the issue could be avoided by conducting a careful causal analysis.

<sup>5</sup>In causality literature, subjects of a study – e.g., patients in a clinical trial – are called units. In our case, each flight is considered a unit.

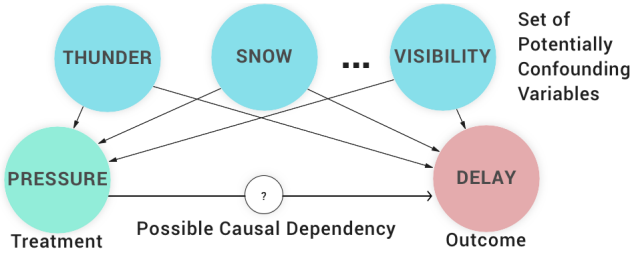


Figure 4: Confounding influence

causal link between a treatment and outcome. This is often the case in real world analysis as the myriad of variables involved greatly complicate understanding a phenomenon. In this example, we will show that the observed positive effect of LowPressure was actually the result of the confounding influence of other factors including low visibility, snow, and thunder. Specifically, we will show that Low Pressure is highly associated with unsettled weather conditions such as LowVisibility (see Fig. 4). The average of LowVisibility in the treated group is much less than that of the opposite group. Thus, it is unclear that the observed DepDelay difference between the groups is caused by LowPressure or LowVisibility.

**Adjusting for confounding variables:** For the sake of drawing valid conclusions to the research questions, we must adjust for these confounding variables. Conceptually, adjusting for one confounding variable is easy. It involves partitioning the data into similar groups with similar confounding influence measures. Next, the ATE is computed as the weighted average of the effects *in each group*. However, real world causal inference is much more complicated as it involves many confounding variables. Fig. 4 shows some of the confounding variables in this example. In this case, the groups become overly specific and begin to lack enough units to create a meaningful calculation of the ATE.

Sophisticated techniques are required to adjust for a large set of confounding variables. ZaliQL implements two primary approaches that are commonly found in social sciences and statistics: *coarsened exact matching (CEM)* and *propensity score matching (PSM)* [5, 10]. The goal of matching is to cut data so that the remaining *matched* data have greater *balance* between the treatment and control groups. In other words, the empirical distributions of the confounding variables in the treated and control groups are similar after matching. Once the groups have more or less achieved a sense of balance, the observed difference of the outcome between the two groups can be attributed to the treatment. In our demonstration, we will select a set of covariates deemed to confound the treatment of DepDelay (Fig. 3c) and we will select a matching method and adjust its tuning parameters.

**Checking Balance:** This next step requires verifying that the matching process improved the covariates balance. Specifically, we will compare the distribution for each covariate between the treated and control group on the matched data for all treatments. ZaliQL provides both numerical and visual summaries for this step of the process including quartile-quartile and mean difference plots. This can be seen in Fig. 3d and Fig. 3e.

**Answers to our causal questions:** Finally, we are able to answer the causal questions created in the first step of our process. For Q1, we will show that LowPressure has no significant causal effect on DepDelay. For Q2, we will

Unit	Covariates $X$	Treatment assignment $T$	Treatment outcome $Y(1)$	Control outcome $Y(0)$	Causal Effect $Y(1) - Y(0)$
1	$X_1$	$T_1$	$Y_1(1)$	$Y_1(0)$	$Y_1(1) - Y_1(0)$
2	$X_2$	$T_2$	$Y_2(1)$	$Y_2(0)$	$Y_2(1) - Y_2(0)$
...	...	...	...	...	...
$N$	$X_N$	$T_N$	$Y_N(1)$	$Y_N(0)$	$Y_N(1) - Y_N(0)$

Figure 5: The Potential Outcome Framework

identify that other treatments have significant causal effects on DepDelay (Fig. 3f). Regarding Q3, we will report that the major causes of flight delay at the airports included in the study are actually different. T-tests are used to validate the statistical significance of our findings. We also show that the obtained results are validated by FAA reports.

**Scalability:** The last part of this demonstration will allow us to show the scalability by letting users interactively run queries of this large dataset. This process would take existing toolkits hours to complete; however, as seen in Fig. 3g, ZaliQL can process 100M entries in less than a few minutes, whereas other systems such as R, SPSS, or SAS, take up to an hour just to process 5M entries.<sup>6</sup>

## 4. INTERNALS

### 4.1 Basic Formalism

The basic causal model in statistics is called the *potential outcome framework (POF)* [11]. In this model, we are given a table with  $N$  rows called *units* indexed by  $i = 1 \dots N$  (see Table 5). The binary attribute  $T$  denotes *treatment assignment*.  $X$  is a vector of background characteristics, (e.g., airport, airline, weather ...) of each unit, called *covariates*, unaffected by treatment; and the two attributes  $Y(0), Y(1)$  represent *potential outcomes*:  $Y(1)$  is the outcome of the unit if it is exposed to the treatment and  $Y(0)$  is the outcome when it is exposed to the control.

The *treatment effect* caused by the treatment  $T_i$  for the  $i$ th unit is defined as  $Y_i(1) - Y_i(0)$ . The goal of causal analysis is to compute the *average treatment effect (ATE)*:

$$ATE = \mathbb{E}[Y(1) - Y(0)] = \mathbb{E}[Y(1)] - \mathbb{E}[Y(0)].$$

The so-called *fundamental problem of causal inference* is that for each unit we only know either  $Y(1)$  or  $Y(0)$  but not both. For example, each individual flight has either LowPressure=1 or LowPressure=0, so only one of DepDelay(1) or DepDelay(0) is available for each row. Thus, further assumptions are needed for estimating ATE.

The strongest is the *independence assumption*: the treatment mechanism is independent of the potential outcomes, i.e.,  $(Y(1), Y(0)) \perp\!\!\!\perp T$ . This might hold in a properly constructed randomized trial. Then,  $\mathbb{E}[Y(1)] = \mathbb{E}[Y(1)|T = 1]$  and similarly  $\mathbb{E}[Y(0)] = \mathbb{E}[Y(0)|T = 0]$  and so:

$$ATE = \mathbb{E}[Y(1)|T = 1] - \mathbb{E}[Y(0)|T = 0].$$

However, ZaliQL is designed for drawing causal inference from *observational data*, where independence fails in general. Here, statistical literature makes two standard, weaker assumptions called *Strong Ignorability*: for all  $x$ :

- (1)  $Y(0), Y(1) \perp\!\!\!\perp T | X = x$  (unconfoundedness) and
- (2)  $0 < \Pr(T = 1 | X = x) < 1$  (overlap) [10].

If strong ignorability holds, one can estimate ATE by taking the average difference for each group with a particular

<sup>6</sup>We note that the developers of statistical software packages for CEM have identified ZaliQL as a more scalable approach: see <http://gking.harvard.edu/cem>.

value of  $X$  i.e.,  $\mathbb{E}_x[\mathbb{E}[Y(1) - Y(0)|X = x]]$ . In practice, however, a direct application of this method is impossible, because the data is typically very sparse: for any value  $X = x$  we either have no data values at all, or very few such values, which means that estimating  $\mathbb{E}[Y(T)|T = 1 \text{ or } 0, X = x]$  as an average from the database leads to large sampling error. A solution adopted in statistics is *matching* [10].

## 4.2 Matching Methods

This process involves theoretically matching each treated unit to one of multiple control units with similar values of the covariate attributes. Closeness is defined using some distance function between the covariate values of two units. Given a table,  $R(T, X_1 \dots, X_n, Y)$ , ZaliQL offers the following two matching methods.

**Propensity score matching:** The most common method is  $k : 1$  nearest neighbor matching based on *propensity score matching (PSM)* [10]. A propensity score is the probability of a unit being assigned to a particular treatment given a set of covariates (i.e.  $P(x) = P(T = 1|X = x)$ ). This method selects the  $k$  best control matches for each individual in the treatment group which are closer than a pre-specified *caliper*. ZaliQL estimates propensity score using logistic regression.

The basic SQL statement to perform PSM is depicted in Fig. 6. In this solution, nearest control units are identified by means of an anti-join. Specifically, all potential matches, and their distances, are identified by joining the treated with the control units that are closer on propensity score than the caliper. Then, this set is sorted into ascending order of distances. In addition, the order of each row in the sorted set is identified using the window function `ROW_NUMBER`. Finally,  $k$  closest controls are selected as the matched units. Note that ZaliQL generates more efficient SQL statements by leveraging recent developments in spatial-databases (see e.g., [7]). The basic SQL statement to perform PSM is depicted in Fig. 6. In this solution, nearest control units are identified by means of an anti-join. In other words, all potential matches, and their distances, are identified by joining the treated with the control units that are closer on propensity score than the caliper. Then, this set is sorted into ascending order of distances. Additionally, the order of each row in the sorted set is identified using the window function `ROW_NUMBER`. Finally,  $k$  closest controls are selected as the matched units. Note that ZaliQL generates more efficient SQL statements by leveraging recent developments in spatial-databases (see e.g., [7]).

**Coarsened exact matching (CEM):** In this method, the vector of covariates  $X$  is coarsened according to a set of user-defined cut points or any automatic discretization algorithm. All units with similar coarsened covariate values are placed in unique groups. These unique groups with at least one treated and one control unit are retained while all others are discarded from the remainder of the process. We let  $X$  be the coarsened version of  $X$  and  $R^c$  be a coarsened version of  $R$ . As depicted in Fig. 7, CEM is essentially a GROUP-BY-HAVING query. ZaliQL also supports several supervised and unsupervised methods for coarsening that have been proposed [1].

We observed that CEM is an “Iceberg query” in the sense that it is a GROUP-BY-HAVING query where the output size is typically much smaller than the input (i.e. the tip of an iceberg). Iceberg queries have been studied extensively

```
CREATE VIEW PSM_Matched
AS WITH potential_matches AS
  (SELECT treated.ID AS tID, control.ID AS cID,
    abs(Treated.PS-Control.PS) AS distance
  FROM R AS control, R AS treated
  WHERE control.T=0 AND treated.T=1
    AND abs(Treated.PS-Control.PS) < caliper),
  ordered_potential_matches AS
  (SELECT *, ROW_NUMBER() over (ORDER BY distance) AS order
  FROM potential_matches)
SELECT *
FROM ordered_potential_matches AS rp
WHERE NOT EXISTS
  (SELECT *
  FROM ordered_potential_matches AS z
  WHERE z.order < opm.order AND z.cID=opm.cID)
AND (SELECT count(*)
  FROM ordered_potential_matches AS opm
  WHERE z.order < opm.order AND z.tID=opm.tID) ≤ k;
```

Figure 6: Propensity score matching

```
CREATE VIEW CEM_Matched AS
WITH subclasses AS
  (SELECT *,
    max(ID) OVER w subclass, max(T) AS min_treatment,
    min(T) AS max_treatment
  FROM Rc
  Group by  $\mathcal{X}$ 
  Having min_treatment!=max_treatment)
SELECT *
FROM subclasses, Rc
WHERE subclasses. $\mathcal{X}$ =Rc. $\mathcal{X}$ 
```

Figure 7: Coarsened Exact Matching

in databases and data mining [2, 3]. ZaliQL leverages these techniques and the prior research on them in order to more efficiently compute CEM for several treatments simultaneously, giving it a strong advantage over other softwares.

**Analysis after matching:** After a balanced matched subset of data is extracted, ATE can be computed. ZaliQL supports a wide range of statistical tests such as z-test, t-test, linear regression t-test, and Chi-Square test to compute the statistical significance of the treatment effect. After a balanced and matched subset of data is extracted, ATE can be computed. ZaliQL supports a wide range of statistical tests such as z-test, t-test, linear regression t-test, and Chi-square test to compute the statistical significant of the treatment group.

## 5. CONCLUSIONS

In this demonstration, we will introduce ZaliQL, a tool for performing causal inference on large relational data within a DBMS. ZaliQL makes the first step towards truly scalable causal inference by modeling it as a data management problem. ZaliQL implements a wide range of methods for causal inference developed in statistics with existing techniques in data management. This provides scalable evaluation of several causal hypothesis on relational data. Overall, this demonstration will serve as both an introduction to causal inference and an illustration of the vast benefits ZaliQL possesses over traditional statistical analysis packages.

## 6. ACKNOWLEDGEMENTS

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