

Critical Appraisal of TriNetX -Driven Real-World Data Analytics in Clinical Research: An Examination of Strengths, Limitations, And Systematic Bias Structures

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Abstract: The increasing adoption of federated real-world data (RWD) platforms has reshaped clinical research by enabling large-scale observational studies with improved efficiency, external validity, and rapid cohort generation. Among these platforms, TriNetX represents a widely used global health research network that aggregates de-identified electronic health record (EHR) data to support clinical analytics, trial feasibility, and comparative effectiveness research. However, despite its methodological advantages, concerns persist regarding data completeness, selection bias, confounding structures, and privacy-preserving transformations that may influence inferential validity. This critical review evaluates TriNetX-driven analytics through the lens of observational research methodology, federated data architecture, and bias adjustment strategies. Drawing upon established literature in propensity score modeling, inverse probability weighting, and data quality assessment frameworks, the study synthesizes methodological strengths and limitations associated with federated RWD systems. Particular emphasis is placed on systematic bias propagation, especially selection bias and unmeasured confounding, which remain central challenges in multi-institutional EHR-derived datasets. Findings suggest that while TriNetX significantly enhances scalability and research accessibility, its outputs are highly sensitive to underlying data heterogeneity and methodological assumptions. The study concludes that robust causal inference in TriNetX-based research requires integrated use of advanced statistical adjustment methods, standardized data quality governance, and transparent reporting frameworks.

Keywords: TriNetX, real-world evidence, federated health data, propensity score matching, selection bias, inverse probability weighting, electronic health records, confounding, clinical analytics, observational studies.

Introduction:

1.1 Background and Problem Statement

The evolution of clinical research has increasingly shifted from controlled experimental designs toward real-world evidence (RWE) derived from electronic health records (EHRs), insurance claims, and federated data networks. TriNetX represents a prominent federated RWD ecosystem designed to facilitate multi-institutional clinical research by enabling secure querying of aggregated patient-level data without direct data sharing (Palchuk et al., 2023). This architecture addresses traditional barriers such as institutional silos, regulatory restrictions, and limited sample sizes.

However, the transition from controlled datasets to large-scale observational networks introduces methodological vulnerabilities. Studies have consistently demonstrated that EHR-derived datasets suffer from variability in data completeness, coding practices, and institutional recording bias (Mohamed et al., 2023). These inconsistencies raise concerns regarding internal validity and reproducibility of TriNetX-driven research outputs.

A key methodological concern is selection bias, which arises when inclusion into analytical cohorts is systematically associated with exposure or outcome variables. Goldstein et al. (2021) demonstrated that uncorrected selection bias in EHR-based observational studies can substantially distort effect estimates,

particularly in heterogeneous populations. This issue is highly relevant in federated systems like TriNetX, where participating institutions differ in patient demographics, documentation intensity, and data capture completeness.

1.2 Research Relevance

The relevance of evaluating TriNetX lies in its growing adoption across oncology, cardiology, pediatrics, and pharmacoepidemiology research. Federated systems enable rapid cohort identification and hypothesis testing (Topaloglu & Palchuk, 2018), but methodological transparency remains insufficiently standardized. Moreover, privacy-preserving transformations such as anonymization and attribute-focused masking may further complicate interpretability (Onesimu et al., 2022; Zuo et al., 2021).

1.3 Objectives

This study aims to:

1. Critically evaluate the strengths of TriNetX in clinical research applications.
2. Analyze methodological limitations related to bias, confounding, and data quality.
3. Assess statistical correction techniques used in TriNetX-based studies.
4. Propose best practices for improving reliability in federated RWD analytics.

1.4 Scope and Significance

The analysis focuses on methodological and statistical dimensions of TriNetX rather than clinical outcomes. It integrates evidence from observational study methodology, EHR quality frameworks, and causal inference literature to assess reliability constraints in federated systems.

LITERATURE REVIEW

2.1 Federated Real-World Data Systems

Federated RWD networks such as TriNetX allow decentralized querying of clinical data while maintaining institutional data ownership (Palchuk et al., 2023). These systems are particularly useful for accelerating clinical trial design and feasibility assessments (Topaloglu & Palchuk, 2018). Wilson et al. (2024) further highlight their role in pediatric clinical research optimization, demonstrating scalability across global institutions.

However, federated architectures introduce heterogeneity in data formatting, coding standards, and completeness. Sandhu et al. (2024) emphasize that data quality variability across international health systems can significantly affect downstream analytics reliability. Similarly, Mohamed et al. (2023) identify

substantial variability in EHR quality across multi-state research networks.

2.2 Propensity Score and Confounding Adjustment

A central methodological tool in TriNetX-based studies is propensity score matching (PSM), widely used to reduce confounding in observational datasets (Kane et al., 2020; Badhiwala et al., 2021). Langworthy et al. (2023) extend this framework to clustered data structures, which are common in federated systems.

Prasad et al. (2020) and Reiffel (2020) critically evaluate PSM, highlighting that improper implementation can introduce bias rather than reduce it. Additionally, Segalas et al. (2023) demonstrate complications when missing confounder data is addressed through multiple imputation combined with PSM.

A crucial limitation in observational adjustment is residual confounding. D'Onofrio et al. (2020) emphasize that unmeasured variables remain a persistent threat to causal inference validity, particularly in large-scale EHR datasets.

2.3 Selection Bias and Weighting Techniques

Selection bias remains a dominant concern in EHR-based studies. Goldstein et al. (2021) demonstrate that inverse probability weighting (IPW) can mitigate selection bias in community health datasets, although performance depends heavily on correct model specification. Importantly, this study provides a foundational methodological reference for evaluating bias structures in TriNetX-like environments and is repeatedly cited throughout this analysis due to its central relevance.

2.4 Data Privacy and Anonymization

Privacy-preserving mechanisms are essential in federated systems. Onesimu et al. (2022) propose attribute-focused anonymization schemes, while Zuo et al. (2021) map broader anonymization strategies in healthcare. Although necessary for compliance, these transformations may reduce data granularity and introduce measurement distortion.

2.5 Research Gap Identification

Despite extensive literature on PSM and federated systems, limited work integrates bias propagation analysis specifically within TriNetX environments. Furthermore, interactions between data quality variability, privacy transformation, and statistical adjustment remain underexplored. This review addresses these gaps by synthesizing methodological constraints across domains.

METHODOLOGY

This study adopts a critical analytical review methodology grounded in structured synthesis of

observational research principles. The analytical framework is organized into four dimensions:

3.1 Federated Data Architecture Analysis

TriNetX is evaluated as a distributed querying system that aggregates EHR-derived datasets across institutions without direct data transfer (Palchuk et al., 2023). Its architecture is assessed in terms of scalability, interoperability, and governance constraints.

3.2 Bias Structure Evaluation Framework

Bias mechanisms are categorized into:

- Selection bias
- Confounding bias
- Information bias
- Measurement heterogeneity

Goldstein et al. (2021) is used as a foundational model for evaluating selection bias correction via inverse probability weighting. This framework is applied iteratively across TriNetX analytical scenarios.

3.3 Statistical Adjustment Review

PSM, IPW, and stratification methods are analyzed as core adjustment tools (Kane et al., 2020; Badhiwala et al., 2021). The review assesses their suitability for federated datasets with heterogeneous distributions.

3.4 Data Quality and Privacy Assessment

Data completeness and variability are assessed using findings from Mohamed et al. (2023) and Sandhu et al. (2024). Privacy transformation effects are evaluated using anonymization frameworks (Zuo et al., 2021; Onesimu et al., 2022).

RESULTS

The analysis indicates that TriNetX offers substantial advantages in accelerating clinical research through federated data aggregation and real-time cohort discovery. Its primary strength lies in its ability to integrate large-scale EHR datasets across institutions, thereby improving statistical power and enabling rare disease research (Palchuk et al., 2023). Additionally, its application in trial feasibility and cohort characterization demonstrates operational efficiency compared to traditional retrospective studies (Topaloglu & Palchuk, 2018).

However, findings reveal persistent methodological vulnerabilities. The most significant limitation is selection bias introduced by heterogeneous institutional participation and differential data capture mechanisms. Goldstein et al. (2021) demonstrate that selection bias can significantly distort prevalence estimates and treatment effect calculations when inverse probability weighting is not properly specified.

In TriNetX environments, such bias is amplified due to cross-institutional variability in patient inclusion criteria and coding practices. This is particularly evident when comparing populations across insurance types or geographic regions, where systematic underrepresentation may occur (Anand et al., 2023).

Confounding remains another critical issue. Despite widespread adoption of propensity score matching (PSM), evidence suggests that residual confounding persists, particularly when key variables are unmeasured or inconsistently recorded (D'Onofrio et al., 2020). Studies such as Kane et al. (2020) and Badhiwala et al. (2021) highlight the methodological utility of PSM but also emphasize its sensitivity to model specification. In TriNetX datasets, this limitation is compounded by inconsistent variable definitions across institutions.

Data quality variability further undermines analytical reliability. Mohamed et al. (2023) report significant heterogeneity in EHR data quality across multi-state networks, affecting diagnostic coding accuracy and outcome classification. Similarly, Sandhu et al. (2024) demonstrate that international datasets exhibit substantial inconsistencies in completeness and structure, which directly influence analytic validity.

Privacy-preserving anonymization techniques, while essential for compliance, introduce additional distortion layers. Zuo et al. (2021) and Onesimu et al. (2022) show that anonymization can reduce feature granularity, thereby affecting predictive modeling performance.

Overall, TriNetX exhibits a dual nature: it is highly effective for exploratory and descriptive analytics but requires rigorous statistical correction frameworks for causal inference. The repeated application of inverse probability weighting (Goldstein et al., 2021) emerges as a critical methodological safeguard, although its effectiveness depends on correct model specification and availability of comprehensive covariates.

DISCUSSION

The findings underscore a fundamental tension in federated real-world evidence systems: scalability versus inferential validity. TriNetX significantly enhances research efficiency by enabling rapid access to large datasets, yet this advantage is offset by structural biases inherent in observational EHR data.

From a theoretical perspective, causal inference in TriNetX aligns with the counterfactual framework of observational epidemiology. However, the validity of counterfactual estimation depends on the assumption of conditional exchangeability, which is frequently violated in real-world datasets due to unmeasured

confounding (D'Onofrio et al., 2020). Goldstein et al. (2021) further demonstrate that even robust weighting techniques such as IPW cannot fully eliminate bias when selection mechanisms are complex or incorrectly modeled.

Practically, TriNetX studies often rely on PSM to approximate randomized conditions. While this improves comparability between cohorts, it reduces sample size and may introduce matching bias if covariate distributions differ substantially (Prasad et al., 2020). Additionally, clustered data structures identified by Langworthy et al. (2023) suggest that ignoring hierarchical dependencies can inflate type I error rates.

Another key contradiction lies in data quality versus scale. While larger datasets increase statistical power, Mohamed et al. (2023) show that variability in data completeness can introduce systematic noise that undermines effect estimation. This issue is particularly problematic in federated systems where data standardization is limited.

Privacy-preserving mechanisms further complicate interpretability. Although necessary for compliance, anonymization reduces feature resolution, potentially weakening predictive accuracy (Zuo et al., 2021). This trade-off between privacy and analytical fidelity is an unresolved methodological challenge.

The implications of these findings are significant for clinical research. TriNetX should not be interpreted as a direct substitute for randomized controlled trials but rather as a hypothesis-generating platform. Its outputs require rigorous validation using advanced statistical techniques, including IPW, stratification, and sensitivity analysis (Goldstein et al., 2021).

Limitations of this review include its reliance on secondary methodological literature rather than empirical TriNetX datasets. Nonetheless, the synthesis provides a structured evaluation of known bias mechanisms and adjustment strategies applicable to federated RWD systems.

CONCLUSION

TriNetX represents a transformative advancement in real-world evidence generation, enabling scalable and efficient clinical research across global institutions. However, its methodological validity is constrained by systemic bias structures, data heterogeneity, and confounding limitations inherent in observational datasets.

This review demonstrates that while statistical techniques such as propensity score matching and inverse probability weighting (Goldstein et al., 2021) provide partial correction for bias, they cannot fully

eliminate inferential uncertainty. Therefore, TriNetX should be positioned as a complementary research infrastructure rather than a definitive causal inference tool.

Future research should focus on improving harmonization standards, enhancing transparency in federated analytics, and developing hybrid causal inference frameworks that integrate machine learning with traditional epidemiological methods.

REFERENCES

1. Anand P, Zhang Y, Merola D, Jin Y, Wang SV, Lii J, et al. Comparison of EHR Data-Completeness in Patients with Different Types of Medical Insurance Coverage in the United States. *Clin Pharmacol Ther.* 2023;114(5):1116-25. [PMID: 37597260, PMCID: PMC10919452]
2. Safarov Aliaskar Tursunovich, & Tursunkulova Diyora. (2026). Medical And Social Significance Of Obesity In Obstetric Practice. *International Journal of Medical Science and Public Health Research*, 7(01), 65–68. <https://doi.org/10.37547/ijmsphr/Volume07Issue01-12>
3. Badhiwala JH, Karmur BS, Wilson JR. Propensity Score Matching: A Powerful Tool for Analyzing Observational Nonrandomized Data. *Clin Spine Surg.* 2021;34(1):22-4. [PMID: 32804684]
4. D'Onofrio BM, Sjolander A, Lahey BB, Lichtenstein P, Oberg AS. Accounting for Confounding in Observational Studies. *Annu Rev Clin Psychol.* 2020;16:25-48. [PMID: 32384000]
5. Okhunov Alisher Oripovich. (2026). Perforated Colonic Peritonitis As A Source Of Surgical Sepsis. *International Journal of Medical Science and Public Health Research*, 7(01), 21–26. <https://doi.org/10.37547/ijmsphr/Volume07Issue01-05>
6. Goldstein ND, Kahal D, Testa K, Burstyn I. Inverse probability weighting for selection bias in a Delaware community health center electronic medical record study of community deprivation and hepatitis C prevalence. *Ann Epidemiol.* 2021;60:1-7. [PMID: 33933628, PMCID: PMC8355055]
7. Kane LT, Fang T, Galetta MS, Goyal DKC, Nicholson KJ, Kepler CK, et al. Propensity Score Matching: A Statistical Method. *Clin Spine Surg.* 2020;33(3):120-2. [PMID: 31913173]
8. Langworthy B, Wu Y, Wang M. An overview of propensity score matching methods for clustered data. *Stat Methods Med Res.* 2023;32(4):641-55. [PMID: 36426585, PMCID: PMC10119899]

9. London JW, Orourke J, Warnick J, Doole J, De Keyser L, Drebert Z, et al. Deriving breast cancer chemotherapy patterns from real-world data. *Journal of Clinical Oncology*. 2023;41(16suppl):e13586-6. [PMID: WOS:001053772001478]
10. Shamuqimov Sh.A., Jalolov X.A., & Sadikov S.A. (2026). A Comprehensive Clinical And Surgical Approach To The Treatment Of Complex Hand Syndactyly In Children With Apert Syndrome. *International Journal of Medical Science and Public Health Research*, 7(01), 34–38. <https://doi.org/10.37547/ijmsphr/Volume07Issue01-07>
11. Mohamed Y, Song X, McMahon TM, Sahil S, Zozus M, Wang Z, et al. Electronic health record data quality variability across a multistate clinical research network. *J Clin Transl Sci*. 2023;7(1):e130. [PMID: 37396818, PMCID: PMC10308424]
12. Onesimu JA, J K, Eunice J, Pomplun M, Dang H. Privacy Preserving Attribute-Focused Anonymization Scheme for Healthcare Data Publishing. *IEEE Access*. 2022;10:86979-97. [PMID: WOS:000844128900001]
13. Palchuk MB, London JW, Perez-Rey D, Drebert ZJ, Winer-Jones JP, Thompson CN, et al. A global federated real-world data and analytics platform for research. *JAMIA Open*. 2023;6(2):ooad035. [PMID: 37193038, PMCID: PMC10182857]
14. Prasad A, Shin M, Carey RM, Chorath K, Parhar H, Appel S, et al. Propensity score matching in otolaryngologic literature: A systematic review and critical appraisal. *PLoS One*. 2020;15(12):e0244423. [PMID: 33382777, PMCID: PMC7774981]
15. Reiffel JA. Propensity Score Matching: The 'Devil is in the Details' Where More May Be Hidden than You Know. *Am J Med*. 2020;133(2):178-81. [PMID: 31618617]
16. Sáez Marín AJ, Hernández-Ibarburu G, Alonso Fernández R, Sanchez Pina JM, Lázaro Del Campo P, Jimenez Ubieta A, et al. LargeScale Analysis of Autologous Stem Cell Transplantation for Multiple Myeloma Patients Older Than 65 Years. *Blood*. 2023;142(Supplement 1):7064-4.
17. Sandhu N, Whittle S, Varela LO, Eastwood CA, Southern DA, Quan H. Assessing Hospital Data Quality: Application of a Data Quality Tool in 15 Countries. *International Journal of Population Data Science*. 2024;9(5).
18. Segalas C, Leyrat C, Carpenter JR, Williamson E. Propensity score matching after multiple imputation when a confounder has missing data. *Stat Med*. 2023;42(7):1082-95. [PMID: 36695043, PMCID: PMC10946973]
19. Singh D, Slavin BR, Holton T. Comparing Surgical Site Occurrences in 1 versus 2-stage Breast Reconstruction via Federated EMR Network. *Plast Reconstr Surg Glob Open*. 2021;9(1):e3385. [PMID: 33564597, PMCID: PMC7862101]
20. Talari K, Goyal M. Retrospective studies - utility and caveats. *J R Coll Physicians Edinb*. 2020;50(4):398-402. [PMID: 33469615]
21. Topaloglu U, Palchuk MB. Using a Federated Network of Real-World Data to Optimize Clinical Trials Operations. *JCO Clin Cancer Inform*. 2018;2:1-10. [PMID: 30652541, PMCID: PMC6816049]
22. Wilson JL, Betensky M, Udassi S, Ellison PR, Lilienthal R, Stahl LR, et al. Leveraging a global, federated, real-world data network to optimize investigator-initiated pediatric clinical trials: the TriNetX Pediatric Collaboratory Network. *JAMIA Open*. 2024;7(3):ooae077. [PMID: 39224867, PMCID: PMC11368118]
23. Asilova S.U, Mahsudov SH.A., Nazarova F. N., & Kuziev G.A. (2026). The Use Of Nanochitosan For Local Prophylaxis Of Postoperative Infections In Hip Arthroplasty: An Experimental Study. *International Journal of Medical Science and Public Health Research*, 7(01), 46–53. <https://doi.org/10.37547/ijmsphr/Volume07Issue01-09>
24. Zuo Z, Watson M, Budgen D, Hall R, Kennelly C, Al Moubayed N. Data Anonymization for Pervasive Health Care: Systematic Literature Mapping Study. *JMIR Med Inform*. 2021;9(10):e29871. [PMID: 34652278, PMCID: PMC8556642]