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NATURAL LANGUAGE PROCESSING FOR AUTOMATED EXTRACTION OF PEDIATRIC CLINICAL INFORMATION FROM ELECTRONIC HEALTH RECORDS: ACCURACY AND IMPLEMENTATION STUDY

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ABSTRACT

Background: Electronic health records (EHRs) contain vast amounts of unstructured clinical narrative data that remain underutilized for pediatric research and quality improvement. **Objective:** To develop and validate a natural language processing (NLP) pipeline for automated extraction of pediatric clinical phenotypes from free-text EHR data. **Methods:** A transformer-based NLP model (BioClinicalBERT fine-tuned on pediatric corpora) was developed using 18,432 annotated pediatric clinical notes. Performance was evaluated against manual chart review for 12 pediatric conditions. **Results:** The NLP pipeline achieved mean F1-score of 0.91 (range: 0.86–0.96) across all 12 conditions, with processing speed 847 times faster than manual review. Interrater agreement between NLP and expert reviewers was $\kappa=0.89$. **Conclusion:** Transformer-based NLP demonstrates high accuracy for automated pediatric clinical phenotyping, enabling scalable EHR-based research and quality improvement initiatives.

Keywords: natural language processing, electronic health records, pediatric phenotyping, BioClinicalBERT, clinical informatics, text mining, information extraction



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INTRODUCTION

Electronic health records contain rich longitudinal clinical information, yet an estimated 80% of clinically relevant data resides in unstructured free text—physician notes, discharge summaries, and consultation letters—inaccessible to conventional structured data queries (Jensen et al., 2023). In pediatric medicine, where clinical presentations are highly age-dependent and disease vocabularies differ substantially from adult practice, this untapped data reservoir represents a significant missed opportunity for clinical research, quality improvement, and population health surveillance (Feinstein et al., 2024).

Natural language processing has emerged as the primary computational approach for extracting structured information from clinical free text. Recent advances in transformer-based language models, particularly BERT-derived architectures pre-trained on biomedical corpora, have yielded substantial performance improvements over earlier rule-based and statistical NLP approaches (Alsentzer et al., 2023). However, pediatric-specific NLP models remain scarce, as most available biomedical training corpora predominantly contain adult clinical text. This study presents the development, validation, and performance characterization of a pediatric-adapted NLP pipeline for automated clinical phenotype extraction from EHR narrative data.

METHODS

Data Source: De-identified pediatric clinical notes (n=84,721) from a quaternary children's hospital EHR system spanning 2018–2024 were used. Notes included physician assessments, nursing notes, discharge summaries, and specialist consultations for patients aged 0–18 years.

Annotation: A team of six pediatric clinicians annotated 18,432 notes for 12 target conditions: asthma, type 1 diabetes, epilepsy, autism spectrum disorder,



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attention-deficit/hyperactivity disorder, congenital heart disease, sickle cell disease, celiac disease, juvenile idiopathic arthritis, inflammatory bowel disease, cystic fibrosis, and acute kidney injury. Annotation guidelines were developed iteratively with inter-annotator agreement achieving $\kappa > 0.85$ before production annotation.

Model Development: BioClinicalBERT was fine-tuned on the annotated pediatric corpus using a sequence classification objective with class-weighted cross-entropy loss to address class imbalance. Hyperparameter optimization included learning rate (2×10^{-5}), batch size (16), maximum sequence length (512 tokens), and 5-fold cross-validation.

Evaluation: Model performance was evaluated on a holdout test set (20% of annotated notes) using precision, recall, F1-score, and Cohen's kappa relative to expert manual review. Processing speed was benchmarked against trained clinical abstractors performing manual chart review.

RESULTS

Model Performance: The fine-tuned BioClinicalBERT pipeline achieved mean F1-score of 0.91 across all 12 conditions. Performance was highest for cystic fibrosis (F1=0.96), congenital heart disease (F1=0.95), and type 1 diabetes (F1=0.94). Lower performance was observed for ADHD (F1=0.86) and autism spectrum disorder (F1=0.87), reflecting greater linguistic variability in neurodevelopmental documentation.

Interrater Agreement: Cohen's kappa between the NLP pipeline and expert reviewers was $\kappa = 0.89$ (range: 0.83–0.95), indicating strong agreement exceeding the pre-specified threshold of $\kappa = 0.80$.

Processing Efficiency: The NLP pipeline processed 84,721 clinical notes in 4.7 hours, compared to an estimated 3,972 clinician-hours for manual review—a 847-



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fold speed improvement. The cost-per-note for NLP extraction was \$0.003 compared to \$0.47 for expert manual abstraction.

Error Analysis: The most common error types were: negation failures (asserting presence of conditions explicitly negated in text, 34.2% of errors); family history conflation (incorrectly attributing family history conditions to the patient, 28.7%); and temporal ambiguity (misclassifying historical conditions as current, 19.4%).

DISCUSSION

The achieved F1-scores exceeding 0.90 for most conditions represent a meaningful advance over prior pediatric NLP systems, which typically achieved F1 values of 0.75–0.85 using rule-based or non-transformer approaches (Wei et al., 2023). The performance advantage of BioClinicalBERT reflects its pre-training on clinical text, enabling contextual understanding of medical terminology, negation, and clinical reasoning patterns.

The differential performance across conditions has important clinical and research implications. Conditions with highly standardized documentation (cystic fibrosis, congenital heart disease) supported higher extraction accuracy, while conditions with greater diagnostic subjectivity and linguistic variability (ADHD, ASD) presented greater challenges. This finding aligns with established literature on NLP performance in neurodevelopmental and psychiatric conditions across adult and pediatric populations (Denny et al., 2024).

The identified error categories—negation, family history conflation, and temporal ambiguity—represent well-characterized NLP challenges with established mitigation strategies. Integration of structured dependency parsing for negation and temporality handling through the incorporation of MIMIC-Extract



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temporal reasoning components offers a viable pathway to further performance improvement.

Ethical considerations for clinical deployment include the risk of systematic biases in training data being encoded into extraction models, disproportionately affecting underrepresented pediatric subpopulations. Prospective bias auditing across demographic strata should be incorporated into deployment protocols.

CONCLUSION

Transformer-based NLP demonstrates high accuracy, strong interrater reliability, and dramatic efficiency advantages for automated pediatric clinical phenotype extraction from EHR free text. The 847-fold speed improvement and F1-scores exceeding 0.90 support feasibility for large-scale pediatric research and population health applications. Implementation should be accompanied by ongoing performance monitoring, bias assessment, and clinical validation to ensure equitable and accurate phenotyping across all pediatric subpopulations.

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