analyzing a pediatric polypharmacy dataset using logistic regression

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1 Introduction

Polypharmacy, often defined as routine use of five or more medications [1], poses a major concern in healthcare, particularly for children. The long-term effects of multiple medications on pediatric development and physiology are not fully understood. Increased rates of chronic conditions like asthma, diabetes, and mental health disorders in children have led to greater medication use, which can increase polypharmacy risks. While sometimes needed for complex or chronic conditions, polypharmacy increases the chance of harmful drug interactions [2], poorer medication adherence [3], and higher healthcare expenses [4]. Identifying factors that predict polypharmacy is essential for creating focused interventions to lessen these risks, especially for vulnerable groups like children.

This project examines the POLYPHARM dataset to determine demographic and clinical predictors of polypharmacy in patients aged 1-19 years. This dataset offers detailed demographic and clinical information, making it a valuable resource for this aim. Given the specific developmental aspects and possible long-term effects of medication regimens in children and adolescents [5], understanding the factors contributing to polypharmacy in this age group is particularly important.

We selected logistic regression because it is appropriate for modeling binary outcomes and aligns with accepted methodological practices [6]. Logistic regression is interpretable, which is valuable for clinical decision-making. It allows for the identification of key predictors and their effect sizes (odds ratios). In contrast to predictive modeling focused on generalizability, this approach allows for the identification of key predictors while providing clinically useful insights. The primary goal is to understand *why* polypharmacy occurs in this population, rather than just predicting its occurrence.

Our analysis found age, gender, and comorbidities to be predictors of polypharmacy. Older age and male gender were associated with higher odds of polypharmacy, while the presence of comorbidities was linked to reduced likelihood. These findings are consistent with prior research on polypharmacy in various populations [2],[9], [10] and provide useful information for healthcare providers managing polypharmacy risks in younger patients.

2 Background

Identifying patterns and predictors of polypharmacy is essential for designing evidence-based interventions that lessen negative risks and improve patient care[3], especially in pediatric

populations where developmental factors require specific strategies [7]. In children, medication regimens can have long-term developmental consequences. Therefore, polypharmacy in this age group presents distinct challenges due to ongoing development and the potential for lasting impacts on growth and overall health [8]. Understanding the factors that contribute to polypharmacy in this age group is vital for improving medication management and patient outcomes.

Prior studies have shown that demographic factors, like age and gender, and clinical characteristics such as comorbidities and healthcare service use, are potential predictors of polypharmacy. For example, older patients and individuals managing multiple chronic conditions are frequently found to have higher medication counts [2]. Gender differences are also relevant, with females often showing higher rates of polypharmacy, possibly due to variations in healthcare-seeking behaviors and prescribing practices [2],[9]. Furthermore, disparities in healthcare access and prescribing patterns across racial and ethnic groups have been reported, further highlighting the need to examine these factors in the context of polypharmacy [10]. Social determinants of health, such as socioeconomic status, healthcare access, and health literacy, are important in shaping medication use and may contribute to differences in polypharmacy rates across different populations. While these factors were not explored in this study, they are important areas for future research to better understand the broader context of polypharmacy in pediatric populations.

This project is based on principles described in "Applied Logistic Regression" by Hosmer et al. (Chapter 3, Section 3.5), which highlights the utility of logistic regression for models involving binary outcomes [6]. Logistic regression was chosen for this study because it is widely understood, interpretable, and well-suited for identifying relationships between predictors and a binary outcome like polypharmacy. Unlike predictive modeling focused on generalizability, this study prioritizes exploring relationships to provide practical insights for healthcare providers.

Using data from the POLYPHARM dataset, which includes 3,500 observations and detailed demographic (e.g., age, sex) and clinical variables (e.g., comorbidities, healthcare utilization), this project applies logistic regression to connect theoretical understanding (using statistical models) with practical applications (in this case, healthcare management). By identifying and quantifying key predictors, the study adds to the growing literature aimed at addressing issues related to polypharmacy.

3 Methods

This section describes the dataset's key features, identifies relationships between variables, and visualizes trends to inform the modeling process.

3.1 Dataset

The POLYPHARM dataset includes 3,500 observations across 14 variables, covering demographic, clinical, and healthcare utilization data. Categorical variables, including POLYPHARMACY and RACE, were converted to factors for appropriate handling during analysis. The variable

types included numeric (such as AGE), binary nominal (such as POLYPHARMACY), multi-level nominal (such as RACE), and ordinal (such as NUMPRIM). Table 1 summarizes the key variable

| Variable | Description |
|--------------|--|
| AGE | Patient age (years) |
| GENDER | Gender (0 - Female, 1 - Male) |
| RACE | Race Categories (0 - White, 1 - Black, 2 - Other) |
| NUMPRIM | Number of primary diagnosis (0 - none, 1 - one, 2 - more than one) |
| COMORBID | Comorbidities (0 - None, 1 - Present) |
| POLYPHARMACY | Polypharmacy status (0 - None, 1 - Present) |

Table 1: Key Variables in the Dataset

The dataset contained one missing value under the URBAN variable and 49 missing values under the NUMPRIM variable. Given the small proportion of missing data (1.4% of observations), we used listwise deletion instead of imputation.

The dataset is publicly available in the aplore3 R package, which includes datasets from Hosmer, Lemeshow, and Sturdivant's "Applied Logistic Regression" (3rd Ed., 2013) [6]. The dataset can be accessed at https://rdrr.io/cran/aplore3/man/polypharm.html.

3.2 Exploratory Data Analysis

The exploratory data analysis (EDA) phase aimed to uncover patterns and relationships within the POLYPHARM dataset, guiding the selection of predictors for the logistic regression model. The dataset revealed the following information:

Age distribution: Patients ranged from 1.17 to 18.92 years with an average age (mean) of 11.65 years. Polypharmacy cases had a higher mean age (12.19 years) compared to nonpolypharmacy cases (11.48 years), showing that older individuals are more likely to experience polypharmacy. Figure 1 suggests that patients with polypharmacy tend to be older than those without.

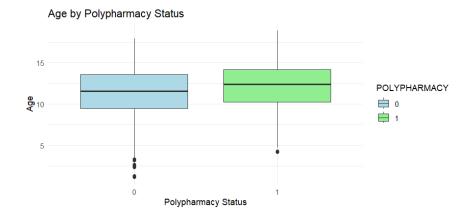


Figure 1: Age Distribution by Polypharmacy Status. All the five boxplot values are higher for the class where polypharmacy is present.

Gender Composition: Female patients accounted for 77.19% of the dataset, reflecting possible differences in healthcare-seeking behavior by gender. Figure 2 shows the polypharmacy outcomes for each gender:

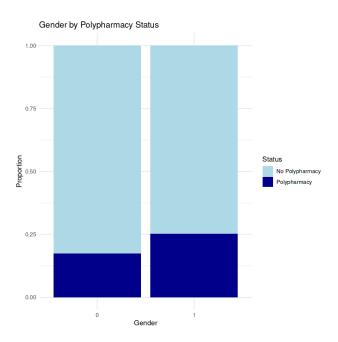


Figure 2: Gender Distribution by Polypharmacy Status

Polypharmacy prevalence: Approximately 23.4% of patients were classified as polypharmacy cases, highlighting its relevance in the dataset.

3.3 Statistical Tests

Pearson's Chi-squared test identified a significant association between RACE and POLYPHARMACY (X-squared = 11.417, df = NA, p = 0.002499). Since this test is the standard method for assessing associations between categorical variables, it follows that we have support for the inclusion of race in the model. A two-sided Fisher's Exact test for count data that gives 0.002386 as p-value also indicates that polypharmacy status varies significantly across racial groups.

3.4 Correlation Analysis

To explore relationships between variables, we computed appropriate measures of correlation for each pair of variable types. The correlation measures for each pair is enumerated in the following table.

| Variable Type Pair | Correlation Measure |
|--|-----------------------|
| Nominal vs. Nominal, Nominal vs. Ordinal, Nominal vs. Binary | Cramér's V |
| Numeric vs. Nominal | Eta-Squared |
| Numeric vs. Numeric | Pearson's Correlation |
| Numeric vs. Binary Nominal | Point-Biserial |
| Numeric vs. Ordinal, Ordinal vs. Ordinal, Binary vs. Ordinal | Spearman's Rank |
| Binary Nominal vs. Binary Nominal | Phi Coefficient |

Table 2: Correlation measures for different variable type pairs.

We saw some several notable relationships:

- AGE and YEAR: A strong positive correlation (r = 0.69) suggesting these variables have a substantial overlap in the directionality of the information they represent.
- NUMPRIM and ANYPRIM: A near perfect correlation (r = 0.99) indicating these variables essentially capture the same information.
- RACE and ETHNIC: A moderate correlation (r = 0.47), suggesting some demographic clustering.

Figure 3 shows the heatmap of the matrix whose entries are the computed correlation measure between the corresponding pair of variables, with darker shades indicating stronger correlations. The heatmap provides a visual summary of the relationships between variables, guiding the selection of predictors for the logistic regression model.

Full Correlation Matrix Heatmap

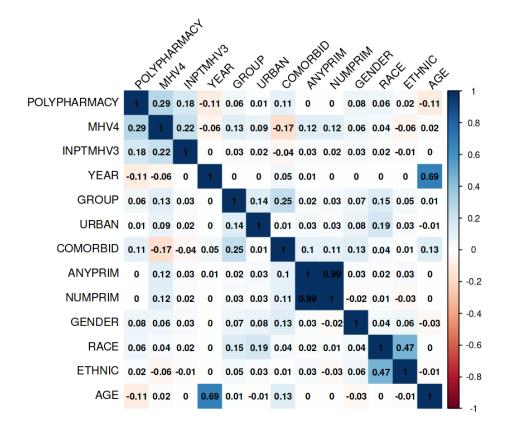


Figure 3: Correlation Matrix

3.5 Logistic Regression Model

The logistic regression model was fitted to the entire dataset to explore the relationships between the selected predictor variables and the likelihood of polypharmacy. The predictor variables included in the model were age (AGE), gender (GENDER), race (RACE, ETHNIC), number of primary diagnosis (NUMPRIM), and presence of comorbidities (COMORBID). These variables were selected based on their clinical relevance, insights from the exploratory data analysis, and prior research [2],[9], [10].

A visual inspection of the boxplot in Figure 1 suggests a potential linear relationship between age and the logit of polypharmacy status. The assumption of a linear relationship between continuous predictors and the logit of the outcome gives us some justification to include AGE as a continuous predictor.

The logistic regression model was specified as

$$logit(P) = \beta_0 + \beta_1 AGE + \beta_2 GENDER + \beta_3 NUMPRIM + \beta_4 COMORBID + \beta_5 RACE + \beta_6 ETHNIC.$$

In the equation, P represents the probability of polypharmacy occurrence, and the β coefficients represent the log odds for each predictor variable.

The logistic regression model was implemented using the 'glm' function in R with a binomial family. After fitting the logistic regression model, we assessed multicollinearity among the predictors used in the model by computing variance inflation factors (VIFs). All VIFs were below 5, indicating no significant multicollinearity among the included predictors and supporting the stability of the logistic regression coefficients.

| Predictor | GVIF | Df | $\mathrm{GVIF}^{1/(2^*\mathrm{Df})}$ |
|-----------|----------|----|--------------------------------------|
| AGE | 1.011251 | 1 | 1.005610 |
| GENDER | 1.014441 | 1 | 1.007195 |
| NUMPRIM | 1.013536 | 2 | 1.003367 |
| COMORBID | 1.028276 | 1 | 1.014039 |
| ETHNIC | 1.206754 | 1 | 1.098524 |
| RACE | 1.206079 | 2 | 1.047958 |

Table 3: R output from Variance Inflation Factor (VIF) computation. GVIF values below 5 indicate no significant multicollinearity.

4 Evaluation

This section presents the results of the fitted logistic regression model and discusses its performance. It is important to note that the model was fitted to the entire dataset to prioritize the exploration of relationships, rather than assessing its generalizability to an independent dataset.

4.1 Logistic Regression Results

Our logistic regression analysis identified the following significant predictors of polypharmacy, with odds ratios and confidence intervals indicating the strength and direction of these associations:

- Age: For every one-year increase in age, the odds of polypharmacy increased by 8%. (Odds Ratio = 1.08, 95% CI: 1.05-1.11, p < 0.001).
- Gender: Males were 1.49 times more likely to experience polypharmacy than females (Odds Ratio = 1.49, 95% CI: 1.21-1.84, $p \approx 0.0015$).
- Comorbidity: Patients with comorbidities had significantly lower odds of polypharmacy (Odds Ratio = 0.518, 95% CI: 0.39-0.66, p < 0.001).
- Race (RACE1): Compared to the reference category for race, patients belonging to RACE1 have around 33% lower odds of having polypharmacy (Odds Ratio = 0.675, 95% CI: 0.52-0.84, p < 0.001).

4.2 Model Fit and Residuals

The model's residual deviance decreased from 3776.9 (null model) to 3673.7, indicating a reasonable fit. To formally assess the overall improvement in model fit by including our predictor variables, we conducted a likelihood ratio test comparing our fitted model to the null model (an intercept-only model). Using the deviance reduction of 103.2 with a corresponding reduction of 8 degrees of freedom (corresponding to the number of predictor variables in our model), a likelihood ratio test yielded a highly significant p-value (p < 0.001). This provides further support that the included predictor variables contribute meaningfully to explaining the variation in polypharmacy status.

However, the residual deviance should be interpreted with caution, as the model was fitted to the entire dataset without cross-validation, which may limit its generalizability to other pediatric populations. Minimal extreme residuals suggest that the model adequately captures variability.

4.3 ROC Curve and AUC

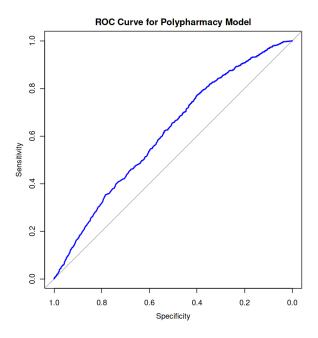


Figure 4: ROC Curve for Polypharmacy Model

The ROC curve (Figure 4) illustrates the trade-off between sensitivity and specificity. The Area Under the Curve (AUC) was 0.611, indicating moderate ability to discriminate between patients who experience polypharmacy and those who do not, within the dataset it was trained on. It is important to interpret this AUC in the context of the model being fitted to the entire dataset, without a separate validation set. While the AUC value suggests that the model can classify polypharmacy cases better than random guessing, there is room for improvement in its classification accuracy. Future studies should consider using advanced techniques such as cross-validation or machine learning algorithms to improve predictive ac-

curacy. These methods could help identify non-linear relationships and interactions between predictors, potentially leading to more accurate risk stratification and targeted interventions.

5 Conclusions and Future Work

This study identified significant demographic and clinical predictors of polypharmacy in a cohort of pediatric patients aged 1-19, providing actionable insights for healthcare providers to manage medication use more effectively in this population. Among the predictors examined, age emerged as the most statistically significant factor in determining polypharmacy risk. However, the presence of comorbidities showed the largest effect size, with these patients being nearly half as likely to experience polypharmacy compared to those without comorbidities. Gender also played a moderate role, with male patients showing a notably higher risk of polypharmacy compared to females.

Specifically, older age, male, absence of comorbidities, and not belonging to RACE1 were associated with a higher likelihood of polypharmacy. These findings highlight key subgroups within the pediatric population who may be at elevated risk of polypharmacy, necessitating closer monitoring and targeted medication management strategies to mitigate adverse outcomes. It is also notable that patients with comorbidities had lower odds of polypharmacy compared to those without comorbidities (Odds Ratio = 0.518, p < 0.001). This suggests that the presence of comorbidities in this dataset was associated with a reduced likelihood of polypharmacy, possibly due to differences in medication management, the nature of the conditions included in the comorbidity variable, or confounding factors such as age and healthcare utilization. The findings provide actionable insights for healthcare providers to manage polypharmacy risks effectively.

The analysis of the logistic regression model provides meaningful insights into polypharmacy risk factors while highlighting important areas for future investigation. While our model demonstrates better-than-random predictive ability (AUC = 0.611), the moderate discriminatory performance suggests that additional factors beyond our current predictors likely influence polypharmacy patterns in pediatric populations. These may include social determinants of health, specific medication classes, and healthcare system factors that were not captured in our dataset.

Several key directions emerge for future research. First, longitudinal studies are needed to assess the long-term developmental impacts of polypharmacy in pediatric populations, particularly examining effects on growth and the risk of chronic conditions in adulthood. Second, investigating healthcare system factors, such as provider prescribing behaviors and access to pediatric specialists, could help explain variations in polypharmacy rates. Third, the role of social determinants of health and specific medication classes should be explored to develop more comprehensive predictive models. Finally, while our logistic regression approach provided valuable insights into key predictors, future studies might benefit from advanced modeling techniques, including machine learning algorithms, to uncover non-linear relationships and improve risk stratification accuracy.

This analysis contributes to the growing body of literature on polypharmacy in pediatric populations, providing valuable, actionable insights for healthcare providers. By understanding the factors associated with polypharmacy, clinicians can implement strategies to optimize

medication use, reduce the risk of adverse events, and improve health outcomes for young patients.

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