# DEVELOPMENT AND PROSPECTIVE VALIDATION OF A PREDICTION MODEL TO IDENTIFY CLINICALLY RELEVANT MEDICATION DISCREPANCIES AT THE EMERGENCY DEPARTMENT

Greet Van De Sijpe<sup>1,2</sup>, Matthias Gijsen<sup>1</sup>, Lorenz Van der Linden<sup>1,2</sup>, Stephanie Strouven<sup>1</sup>, Emily Martens<sup>3</sup>, Nele Persan<sup>3</sup>, Veerle Grootaert<sup>3</sup>, Veerle Foulon<sup>2</sup>, Minne Casteels<sup>2</sup>, Sandra Verelst<sup>1</sup>, Peter Vanbrabant<sup>1</sup>, Sabrina De Winter<sup>1</sup> and Isabel Spriet<sup>1,2</sup>

<sup>1</sup>University Hospitals Leuven, Leuven, Belgium; <sup>2</sup> KU Leuven, Leuven, Belgium; <sup>3</sup>AZ St-Jan Brugge, Brugge, Belgium

## **BACKGROUND AND IMPORTANCE**

Medication discrepancies  $\rightarrow$  avoidable harm

Accurate medication history = essential

Medication reconciliation (MED-REC)

- $\rightarrow$  labor-intensive
- $\rightarrow$  prone to many errors





Prospective multicenter study

- MED-REC at emergency department
- Identifying clinically relevant discrepancies

### Three datasets

- 824 patients  $\rightarrow$  development of model **A**.
- B. 350 patients  $\rightarrow$  temporal validation



4CPS-114



MED-REC due to **limited resources** 

> Need for approach to identify patients at risk

## AIM

Develop and validate a prediction model to identify patients at risk for at least one clinically relevant medication discrepancy at the time of emergency department presentation

"The MED-REC predictor"



RESULTS

#### DEVELOPIV

#### (Figure 1)

119 patients  $\rightarrow$  geographic validation

### **Development and validation**

- Multivariable logistic regression outcome = at least one clinically relevant discrepancy
- **Discrimination and calibration**

probability of discrepancy = linear predictor = log (1-probability of discrepancy)

-1.82 + 0.01\*age - 0.96\*residence(nursing home) - 0.79\*residence(other) -0.09\*n drugs + 0.28\*n ATC A drugs + 0.17\*n ATC C drugs + 0.21\*n ATC N drugs + 1.7\*n ATC P drugs + 0.27\*n ATC R drugs

Probability of discrepancy =  $\frac{1}{1+e^{-linear predictor}}$ 

*Figure 1*. Formula of the prediction model to calculate the probability of having at least one clinically relevant discrepancy



At least one clinically relevant discrepancy observed in 35% (A), 37% (B) and 49% (C)

Final model  $\rightarrow$  8 predictors

#### VALIDATION

# DISCRIMINATION

(Figure 2)

Moderate: AUC 0.66

Retained in all datasets

Better than at random selection

#### **CALIBRATION**

(Figure 3)

Excellent calibration (A and B)

Slight underestimation (C)

*Figure 2*. AUROC curves for the MED-RED predictor in the development (A), temporal validation (B) and geographic validation (C) dataset



## **CONCLUSION AND RELEVANCE**

Prediction model  $\rightarrow$  more **efficient** than selection at random

→ guide rational use of limited resources

Depending on available resources

 Construction of probability threshold and alert
Alert rate to  $\uparrow$  specificity or sensitivity

*Figure 3.* Calibration plots for the MED-RED predictor in the development (A), temporal validation (B) and geographic validation (C) dataset



Model incorporated in electronic health

record

Runs in real-time



- Alerts on structured worklist
- MED-REC of high risk patients by pharmacist or pharmacy technicians