

# Drug Recommendation toward Safe Polypharmacy

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# outline

1. Preamble

2. The Proposed Method

3. Experiments

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# Drug-drug Interaction

## 1. DDIs: Drug-drug interactions

The pharmacological effects of a drug are altered by actions of another drug  
Leading to unpredictable consequences.

## 2. ADRs: Adverse drug reactions

Undesired or harmful reactions due to drug administration.

A recent thread is dedicated **to understanding the interaction patterns among high-order DDIs**, and how such patterns can relate to induced ADRs.

# Two problems

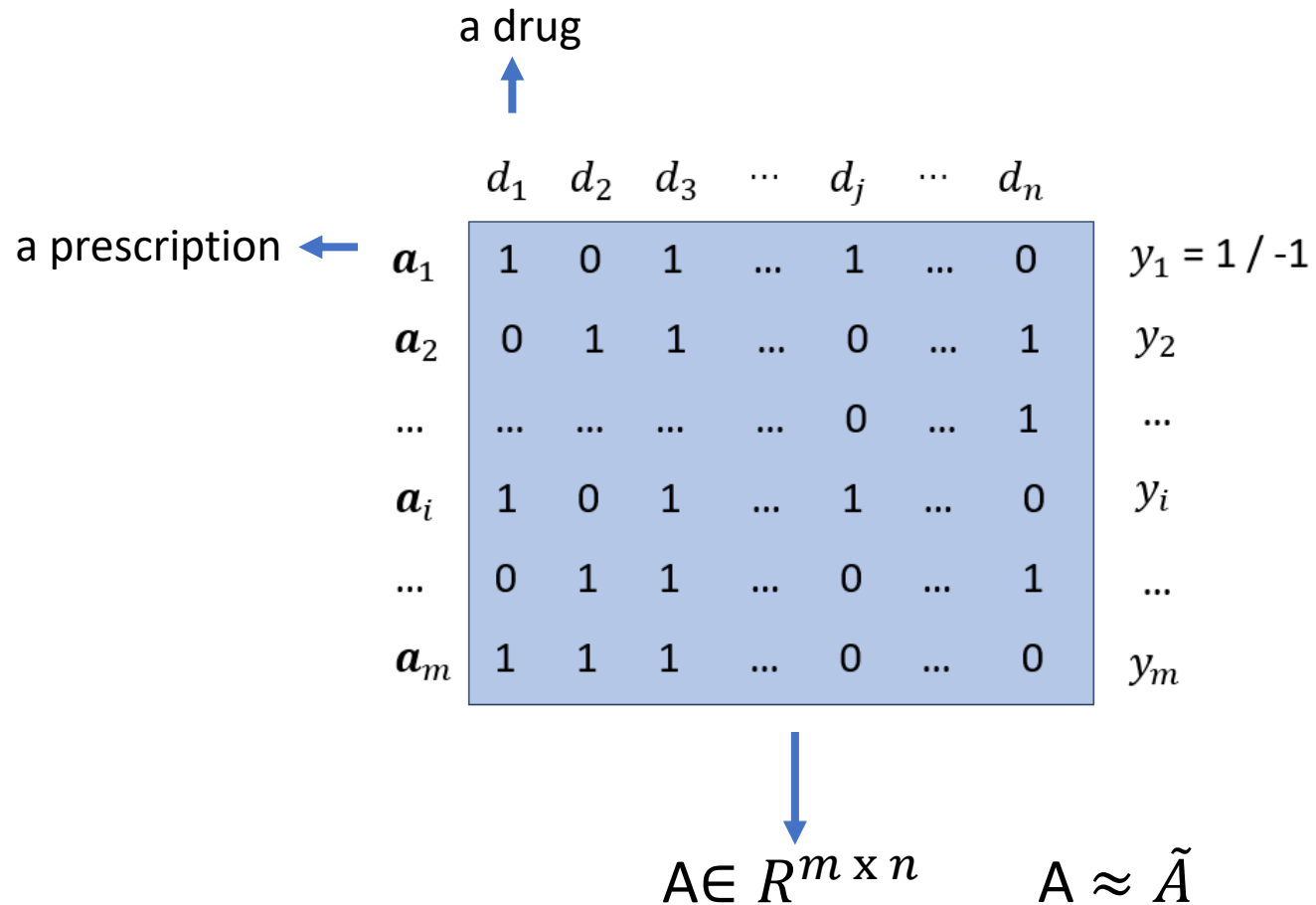
## 1. To-avoid Drug Recommendation:

Given the multiple drugs in a prescription that have been taken simultaneously, **recommend a short of ranked list of drugs** that should be **avoided** taking together with the prescription in order to avoid a particular ADR.

## 2. Safe Drug Recommendation

Given the multiple drugs in a prescription that have been taken simultaneously, **recommend a short ranked list of safe drugs** that, if taken **together with** the prescription, are not likely to induce a particular ADR.

# Notations



$A^+$  prescriptions known to induce ADRs

$A^-$  prescriptions known not to induce ADRs

# Two Methods

## 1. SLIM (Sparse Linear Method)

**Motivation:** A matrix decomposition method that decomposes to obtain a sparse matrix of  $W$  that can be used for feature selection and compression of the data.

$$\tilde{A} = AW$$

## 2. LogR (Logistic Regression)

**Motivation:** binary classification

# SLIM:

SLIM learns **a sparse coefficient matrix** for the items in the system solely from user purchase/rating profiles by solving a regularized optimization problem.

$$\tilde{a}_{ij} = \mathbf{a}_i \mathbf{w}_j^T$$

$\tilde{a}_{ij}$  is the estimated score of  $d_j$  in  $\mathbf{a}_i$   $\mathbf{a}_i \cup \{d_j\}$ .

$$\tilde{A} = AW$$

$\mathbf{w}_j^T$  is a sparse column vector of aggregation coefficients

the sparse coefficient matrix

$$W = [\mathbf{w}_1^T, \mathbf{w}_2^T, \dots, \mathbf{w}_n^T]$$

$$\min_W \quad \text{SLIM}(A; W, \alpha, \lambda) = \frac{1}{2} \|A - AW\|_F^2 + \frac{\alpha}{2} \|W\|_F^2 + \lambda \|W\|_{\ell_1}$$

subject to  $W \geq 0, \text{diag}(W) = 0,$

Control the value of W

$$\tilde{A} = AW$$

Control the sparsity of W



# LogR

the probability of a prescription  $a_i$  inducing the ADR is modeled as follows

$$p(y_i | \mathbf{a}_i; \mathbf{x}, c) = (1 + \exp(-y_i(\mathbf{a}_i \mathbf{x}^\top + c)))^{-1}$$

Where  $\mathbf{x}^\top$  and  $c$  are the parameters. To learn the parameters, LogR solves the following optimization problem,

$$\begin{aligned} \min_{\mathbf{x}, c} \quad & \text{LogR}(\mathbf{y} | A; \mathbf{x}, c, \beta, \gamma) \\ & = \sum_{i=1}^m \log\{1 + \exp[-y_i(\mathbf{a}_i \mathbf{x}^\top + c)]\} + \frac{\beta}{2} \|\mathbf{x}\|_2^2 + \gamma \|\mathbf{x}\|_1 \end{aligned}$$

where  $\mathbf{y} = [y_1; y_2; \dots, y_m]$ ,  $\|\mathbf{x}\|_1 = \sum_{i=1}^n |x_i|$ , and  $\|\mathbf{x}\|_2^2 = \sum_{i=1}^n x_i^2$ .

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# SlimLogR

SlimLogR learns the SLIM and LogR components jointly through solving the following optimization problem

$$\begin{aligned}
 & \min_{W^+, W^-, \mathbf{x}, c} \quad \text{SLIM}(A^+; W^+, \alpha, \lambda) + \text{SLIM}(A^-; W^-, \alpha, \lambda) \\
 & \quad \omega \{ \text{LogR}(\mathbf{y}^+ | \tilde{A}^+ \circ M^+; \mathbf{x}, c, \beta, \gamma) + \text{LogR}(\mathbf{y}^- | \tilde{A}^- \circ M^-; \mathbf{x}, c, \beta, \gamma) \} \xrightarrow[M = \mathbb{I}(A)]{M = \mathbf{1}} \begin{matrix} \boxed{\text{SlimLogR}_{\text{in}}} \\ \boxed{\text{SlimLogR}_{\text{ex}}} \end{matrix} \\
 & \text{subject to} \quad \tilde{A}^+ = A^+ W^+, \tilde{A}^- = A^- W^-, \\
 & \quad W^+ \geq 0, W^- \geq 0, \text{diag}(W^+) = 0, \text{diag}(W^-) = 0
 \end{aligned}$$

where  $\mathbb{I}$  is an indicator function ( $(\mathbb{I})(x) = 0$  if  $x = 0$ , 1 otherwise)

# Training SlimLogR

$$\begin{aligned}
 & \min_{W^+, W^-, \mathbf{x}, c} \quad \text{SLIM}(A^+; W^+, \alpha, \lambda) + \text{SLIM}(A^-; W^-, \alpha, \lambda) \\
 & \quad \omega \{ \text{LogR}(\mathbf{y}^+ | \tilde{A}^+ \circ M^+; \mathbf{x}, c, \beta, \gamma) + \\
 & \quad \quad \text{LogR}(\mathbf{y}^- | \tilde{A}^- \circ M^-; \mathbf{x}, c, \beta, \gamma) \} \\
 & \text{subject to} \quad \tilde{A}^+ = A^+ W^+, \tilde{A}^- = A^- W^-, \\
 & \quad W^+ \geq 0, W^- \geq 0, \text{diag}(W^+) = 0, \text{diag}(W^-) = 0
 \end{aligned}$$

$$\begin{aligned}
 & \min_{W^+, W^-, Z^+, Z^-, \mathbf{x}, c} \quad L(W^+, W^-, Z^+, Z^-, \mathbf{x}, c, \mathbf{u}^+, \mathbf{u}^-, \rho^+, \rho^-) = \\
 & \quad \text{SLIM}(A^+; W^+, \alpha, \lambda) + \text{SLIM}(A^-; W^-, \alpha, \lambda) \\
 & \quad \omega \{ \text{LogR}(\mathbf{y}^+ | \tilde{B}^+ \circ M^+; \mathbf{x}, c, \beta, \gamma) + \\
 & \quad \quad \text{LogR}(\mathbf{y}^- | \tilde{B}^- \circ M^-; \mathbf{x}, c, \beta, \gamma) \} \\
 & \quad \mathbf{u}^{+\top} \mathbf{v}^+ + \frac{\rho^+}{2} \|\mathbf{v}^+\|_2^2 + \mathbf{u}^{-\top} \mathbf{v}^- + \frac{\rho^-}{2} \|\mathbf{v}^-\|_2^2, \\
 & \text{subject to} \quad \tilde{B}^+ = A^+ Z^+, \tilde{B}^- = A^- Z^-, \\
 & \quad \mathbf{v}^+ = \text{vec}(W^+) - \text{vec}(Z^+), \mathbf{v}^- = \text{vec}(W^-) - \text{vec}(Z^-), \\
 & \quad W^+ = Z^+, W^- = Z^-, W^+ \geq 0, W^- \geq 0, \\
 & \quad \text{diag}(W^+) = 0, \text{diag}(W^-) = 0,
 \end{aligned}$$

# Training SlimLogR

$$\begin{aligned}
 & \min_{W^+, W^-, Z^+, Z^-, \mathbf{x}, c, \mathbf{u}^+, \mathbf{u}^-, \rho^+, \rho^-} L(W^+, W^-, Z^+, Z^-, \mathbf{x}, c, \mathbf{u}^+, \mathbf{u}^-, \rho^+, \rho^-) = \\
 & \quad \text{SLIM}(A^+; W^+, \alpha, \lambda) + \text{SLIM}(A^-; W^-, \alpha, \lambda) \\
 & \quad \omega \{ \text{LogR}(\mathbf{y}^+ | \tilde{B}^+ \circ M^+; \mathbf{x}, c, \beta, \gamma) + \\
 & \quad \quad \text{LogR}(\mathbf{y}^- | \tilde{B}^- \circ M^-; \mathbf{x}, c, \beta, \gamma) \} \\
 & \quad \mathbf{u}^{+\top} \mathbf{v}^+ + \frac{\rho^+}{2} \|\mathbf{v}^+\|_2^2 + \mathbf{u}^{-\top} \mathbf{v}^- + \frac{\rho^-}{2} \|\mathbf{v}^-\|_2^2, \\
 & \text{subject to} \quad \tilde{B}^+ = A^+ Z^+, \tilde{B}^- = A^- Z^-, \\
 & \quad \mathbf{v}^+ = \text{vec}(W^+) - \text{vec}(Z^+), \mathbf{v}^- = \text{vec}(W^-) - \text{vec}(Z^-), \\
 & \quad W^+ = Z^+, W^- = Z^-, W^+ \geq 0, W^- \geq 0, \\
 & \quad \text{diag}(W^+) = 0, \text{diag}(W^-) = 0,
 \end{aligned}$$

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## Algorithm 1 Learning SlimLogR

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1: function SlimLogR( $A, \omega, \alpha, \lambda, \beta, \gamma$ )
2:    $\rho^+ = 10, \rho^- = 10, \mathbf{u}_{(0)}^+ = \mathbf{0}, \mathbf{u}_{(0)}^- = \mathbf{0}, k = 0$ 
3:    $Z_{(0)}^+ = W_{(0)}^+, Z_{(0)}^- = W_{(0)}^-$ 
4:   learn  $W_{(0)}^+$  and  $W_{(0)}^-$  from SLIM (Section 6.3)
5:   learn  $\mathbf{x}_{(0)}$  and  $c_{(0)}$  from LogR (Section 6.2)
6:   while not converge do
7:      $\{W_{(k+1)}^+, W_{(k+1)}^-\} := \underset{W^+, W^-}{\text{argmin}} L(W_{(k)}^+, W_{(k)}^-, Z_{(k)}^+, Z_{(k)}^-,$ 
        $\mathbf{x}_{(k)}, c_{(k)}, \mathbf{u}_{(k)}^+, \mathbf{u}_{(k)}^-)$ 
8:      $\{Z_{(k+1)}^+, Z_{(k+1)}^-\} := \underset{Z^+, Z^-}{\text{argmin}} L(W_{(k+1)}^+, W_{(k+1)}^-, Z_{(k)}^+, Z_{(k)}^-,$ 
        $\mathbf{x}_{(k)}, c_{(k)}, \mathbf{u}_{(k)}^+, \mathbf{u}_{(k)}^-)$ 
9:      $\{\mathbf{x}_{(k+1)}, c_{(k+1)}\} := \underset{\mathbf{x}, c}{\text{argmin}} L(W_{(k+1)}^+, W_{(k+1)}^-, Z_{(k+1)}^+, Z_{(k+1)}^-,$ 
        $\mathbf{x}_{(k)}, c_{(k)}, \mathbf{u}_{(k)}^+, \mathbf{u}_{(k)}^-)$ 
10:     $\mathbf{u}_{(k+1)}^+ = \mathbf{u}_{(k)}^+ + \rho^+ (\text{vec}(W_{(k+1)}^+) - Z_{(k+1)}^+)$ 
11:     $\mathbf{u}_{(k+1)}^- = \mathbf{u}_{(k)}^- + \rho^- (\text{vec}(W_{(k+1)}^-) - Z_{(k+1)}^-)$ 
12:     $k = k + 1$ 
13:  end while
14:  return  $W_{(k+1)}^+, W_{(k+1)}^-, Z_{(k+1)}^+, Z_{(k+1)}^-, \mathbf{x}_{(k+1)}$  and  $c_{(k+1)}$ 
15: end function

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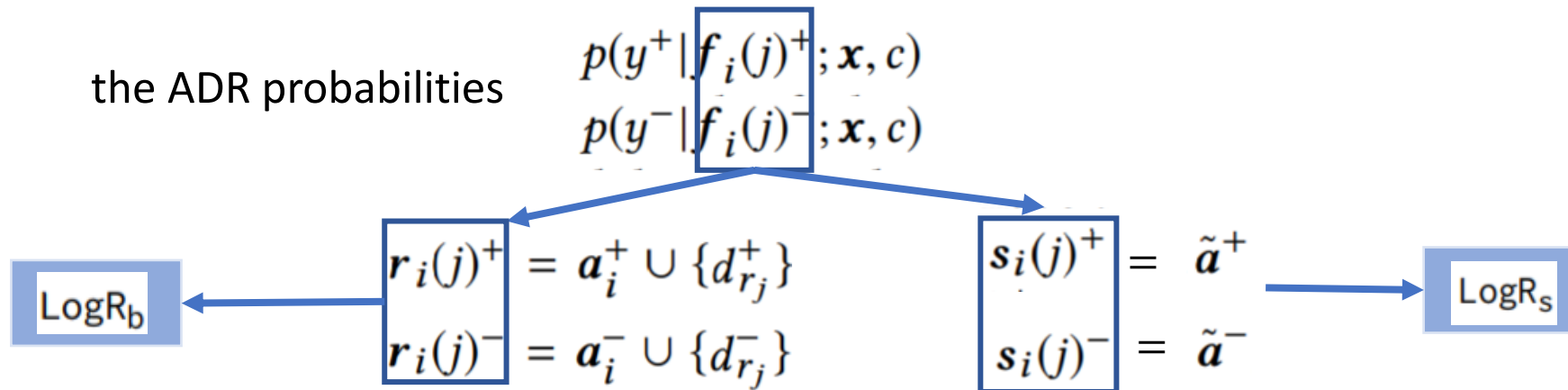
# Applying SlimLogR

- Step 1: use the learned SLIM component to recommend a list of potential to-avoid/safe drug candidates  $\{d_{r_j}\}$ ;

the scores of being potential to-avoid drugs:  $\tilde{a}^+ = aW^+$

the scores of being potential safe drugs:  $\tilde{a}^- = aW^-$

- Step 2: for each recommended candidate  $d_{r_j}$ , combine it with  $a$  (i.e.,  $a \cup \{d_{r_j}\}$ ) and use the learned LogR component to predict the ADR probability of the new prescription.





# Overview

$$\tilde{a}_{ij} = \mathbf{a}_i \mathbf{w}_j^\top$$

$$\min_{W^+, W^-, \mathbf{x}, c}$$

$$\text{SLIM}(A^+; W^+, \alpha, \lambda) + \text{SLIM}(A^-; W^-, \alpha, \lambda)$$

$$\omega \{ \text{LogR}(\mathbf{y}^+ | \tilde{A}^+ \circ M^+; \mathbf{x}, c, \beta, \gamma) +$$

$$\text{LogR}(\mathbf{y}^- | \tilde{A}^- \circ M^-; \mathbf{x}, c, \beta, \gamma) \}$$

SlimLogR<sub>ex</sub>

subject to

$$\tilde{A}^+ = A^+ W^+, \tilde{A}^- = A^- W^-,$$

$$W^+ \geq 0, W^- \geq 0, \text{diag}(W^+) = 0, \text{diag}(W^-) = 0$$

$$p(y_i | \mathbf{a}_i; \mathbf{x}, c) = (1 + \exp(-y_i(\mathbf{a}_i \mathbf{x}^\top + c)))^{-1}$$

the ADR probabilities

$$p(y^+ | f_i(j)^+; \mathbf{x}, c)$$

$$p(y^- | f_i(j)^-; \mathbf{x}, c)$$

$$\mathbf{r}_i(j)^+ = \mathbf{a}_i^+ \cup \{d_{r_j}^+\}$$

$$\mathbf{r}_i(j)^- = \mathbf{a}_i^- \cup \{d_{r_j}^-\}$$

LogR<sub>b</sub>

$$\mathbf{s}_i(j)^+ = \tilde{\mathbf{a}}^+$$

$$\mathbf{s}_i(j)^- = \tilde{\mathbf{a}}^-$$

LogR<sub>s</sub>

$A_{pool}$	mdl	prd	$\overline{\text{rec}}_t$	$\overline{\text{prec}}_t$	$\overline{\text{acc}}_t$
$A_{FAERS}$	SlimLogR <sub>in</sub>	LogR <sub>s</sub>	<b>0.2361</b>	<b>0.2361</b>	<b>0.2958</b>
		LogR <sub>b</sub>	<b>0.2361</b>	<b>0.2361</b>	<b>0.2958</b>
	SlimLogR <sub>ex</sub>	LogR <sub>s</sub>	0.2108	0.2108	0.2795
		LogR <sub>b</sub>	0.2108	0.2108	0.2795
	Rand	-	0.0060	0.0118	0.0238
	LogR	-	0.2082	0.2081	0.2430
	SLIM	-	0.2127	0.2126	0.2811
	SLIM+LogR	-	0.2127	0.2126	0.2813

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# Experiments

Datasets:

dataset	stats	$A^-$		$A^+$	
		$A_-^-$	$A_0^-$	$A_0^+$	$A_+^+$
$A_{FAERS}$	$\#\{a\}$	621,449	1,264	8,986	27,387
	$\#\{d\}$	1,209	417	881	1,201
	avgOrd	6.100	2.351	3.588	7.096
	avgFrq	1.761	225.317	13.730	1.402
	avgOR	-	0.546	16.343	-
$A_*$	$\#\{a\}$	2,200	1,264	2,464	1,000
	$\#\{d\}$	562	417	692	679
	avgOrd	2.678	2.351	3.809	7.615
	avgFrq	42.082	225.317	20.565	5.520
	avgOR	-	0.546	31.998	-

# Experiments

Table 3: Comparison based on the Best  $\overline{rec}_t$  ( $N = 5$ )

$A_{pool}$	mdl	prd	$\overline{rec}_t$	$\overline{prec}_t$	$\overline{acc}_t$
$A_{FAERS}$	SlimLogR <sub>in</sub>	LogR <sub>s</sub>	<b><u>0.2361</u></b>	<b><u>0.2361</u></b>	<b><u>0.2958</u></b>
		LogR <sub>b</sub>	<b><u>0.2361</u></b>	<b><u>0.2361</u></b>	<b><u>0.2958</u></b>
	SlimLogR <sub>ex</sub>	LogR <sub>s</sub>	0.2108	0.2108	0.2795
		LogR <sub>b</sub>	0.2108	0.2108	0.2795
	Rand	-	0.0060	0.0118	0.0238
	LogR	-	0.2082	0.2081	0.2430
	SLIM	-	0.2127	0.2126	0.2811
	SLIM+LogR	-	0.2127	0.2126	0.2813
$A_*$	SlimLogR <sub>in</sub>	LogR <sub>s</sub>	<b><u>0.2618</u></b>	<b><u>0.2615</u></b>	<b><u>0.2815</u></b>
		LogR <sub>b</sub>	<b><u>0.2618</u></b>	<b><u>0.2615</u></b>	<b><u>0.2815</u></b>
	SlimLogR <sub>ex</sub>	LogR <sub>s</sub>	0.2479	0.2479	0.2748
		LogR <sub>b</sub>	0.2479	0.2479	0.2748
	Rand	-	0.0029	0.0059	0.0034
	LogR	-	0.1780	0.1781	0.1032
	SLIM	-	0.2469	0.2467	0.2754
	SLIM+LogR	-	0.2469	0.2467	0.2754

The column "mdl" corresponds to models. The column "prd" corresponds to prediction methods. The best  $\overline{rec}_t$  is underlined. The best overall performance is **bold**.

Table 4: Comparison based on the Best  $\overline{prec}_t$  ( $N = 5$ )

$A_{pool}$	mdl	prd	$\overline{rec}_t$	$\overline{prec}_t$	$\overline{acc}_t$
$A_{FAERS}$	SlimLogR <sub>in</sub>	LogR <sub>s</sub>	<b><u>0.2361</u></b>	<b><u>0.2361</u></b>	<b><u>0.2958</u></b>
		LogR <sub>b</sub>	<b><u>0.2361</u></b>	<b><u>0.2361</u></b>	<b><u>0.2958</u></b>
	SlimLogR <sub>ex</sub>	LogR <sub>s</sub>	0.2108	0.2108	0.2795
		LogR <sub>b</sub>	0.2108	0.2108	0.2795
	Rand	-	0.0060	0.0118	0.0238
	LogR	-	0.2082	0.2081	0.2430
	SLIM	-	0.2127	0.2128	0.2809
	SLIM+LogR	-	0.2127	0.2128	0.2811
$A_*$	SlimLogR <sub>in</sub>	LogR <sub>s</sub>	<b><u>0.2613</u></b>	<b><u>0.2615</u></b>	<b><u>0.2821</u></b>
		LogR <sub>b</sub>	<b><u>0.2613</u></b>	<b><u>0.2615</u></b>	<b><u>0.2821</u></b>
	SlimLogR <sub>ex</sub>	LogR <sub>s</sub>	0.2477	0.2479	0.2759
		LogR <sub>b</sub>	0.2477	0.2479	0.2759
	Rand	-	0.0029	0.0059	0.0034
	LogR	-	0.1780	0.1781	0.1032
	SLIM	-	0.2465	0.2467	0.2754
	SLIM+LogR	-	0.2465	0.2467	0.2754

The column "mdl" corresponds to models. The column "prd" corresponds to prediction methods. The best  $\overline{prec}_t$  is underlined. The best overall performance is **bold**.

Table 5: Comparison based on the Best  $\overline{acc}_t$  ( $N = 5$ )

$A_{pool}$	mdl	prd	$\overline{rec}_t$	$\overline{prec}_t$	$\overline{acc}_t$
$A_{FAERS}$	SlimLogR <sub>in</sub>	LogR <sub>s</sub>	<b><u>0.2355</u></b>	<b><u>0.2355</u></b>	<b><u>0.2969</u></b>
		LogR <sub>b</sub>	<b><u>0.2355</u></b>	<b><u>0.2355</u></b>	<b><u>0.2969</u></b>
	SlimLogR <sub>ex</sub>	LogR <sub>s</sub>	0.2099	0.2099	0.2797
		LogR <sub>b</sub>	0.2108	0.2108	0.2797
	Rand	-	0.0060	0.0118	0.0238
	LogR	-	0.2036	0.2039	0.2468
	SLIM	-	0.2019	0.2020	0.2830
	SLIM+LogR	-	0.2019	0.2020	0.2825
$A_*$	SlimLogR <sub>in</sub>	LogR <sub>s</sub>	<b><u>0.2589</u></b>	<b><u>0.2592</u></b>	<b><u>0.2832</u></b>
		LogR <sub>b</sub>	<b><u>0.2589</u></b>	<b><u>0.2592</u></b>	<b><u>0.2832</u></b>
	SlimLogR <sub>ex</sub>	LogR <sub>s</sub>	0.2477	0.2479	0.2765
		LogR <sub>b</sub>	0.2477	0.2479	0.2765
	Rand	-	0.0029	0.0059	0.0034
	LogR	-	0.1780	0.1781	0.1032
	SLIM	-	0.2434	0.2432	0.2776
	SLIM+LogR	-	0.2434	0.2432	0.2776

The column "mdl" corresponds to models. The column "prd" corresponds to prediction methods. The best  $\overline{acc}_t$  is underlined. The best overall performance is **bold**.

# Parameter Study

$$\begin{aligned}
 &\min_{W^+, W^-, \mathbf{x}, c} \quad \text{SLIM}(A^+; W^+, \alpha, \lambda) + \text{SLIM}(A^-; W^- \alpha \lambda) \\
 &\quad \omega \{ \text{LogR}(\mathbf{y}^+ | \tilde{A}^+ \circ M^+; \mathbf{x}, c, \beta, \gamma) + \\
 &\quad \quad \text{LogR}(\mathbf{y}^- | \tilde{A}^- \circ M^-; \mathbf{x}, c, \beta, \gamma) \} \\
 &\text{subject to} \quad \tilde{A}^+ = A^+ W^+, \tilde{A}^- = A^- W^-, \\
 &\quad W^+ \geq 0, W^- \geq 0, \text{diag}(W^+) = 0, \text{diag}(W^-) = 0
 \end{aligned}$$

**Table 6:**  $\overline{\text{rec}}_t$  of SlimLogR<sub>in</sub> + LogR<sub>s</sub> ( $N = 5$ )
 **Table 7:**  $\overline{\text{prec}}_t$  of SlimLogR<sub>in</sub> + LogR<sub>s</sub> ( $N = 5$ )
 **Table 8:**  $\overline{\text{acc}}_t$  of SlimLogR<sub>in</sub> + LogR<sub>s</sub> ( $N = 5$ )

$\omega \backslash \alpha$	100	50	20	10	5
20	0.2324	0.2344	0.2346	0.2349	0.2338
10	0.2315	0.2352	0.2349	0.2358	0.2344
5	0.2318	0.2355	<b>0.2361</b>	<b>0.2361</b>	0.2341
1	0.2315	0.2327	0.2321	0.2304	0.2290

$A_{pool}$  is  $A_{\text{FAERS}}$ . The best performance is **bold**.

$\omega \backslash \alpha$	100	50	20	10	5
20	0.2325	0.2343	0.2347	0.2349	0.2339
10	0.2315	0.2351	0.2351	0.2359	0.2345
5	0.2321	0.2355	<b>0.2361</b>	0.2359	0.2341
1	0.2315	0.2327	0.2321	0.2305	0.2290

$A_{pool}$  is  $A_{\text{FAERS}}$ . The best performance is **bold**.

$\omega \backslash \alpha$	100	50	20	10	5
20	0.2951	0.2919	0.2890	0.2895	0.2874
10	0.2943	0.2967	0.2938	0.2915	0.2891
5	0.2955	<b>0.2969</b>	0.2958	0.2951	0.2924
1	0.2938	0.2953	0.2936	0.2919	0.2893

$A_{pool}$  is  $A_{\text{FAERS}}$ . The best performance is **bold**.

# Top-N Performance

**Table 9: Top-N Performance of SlimLogR<sub>in</sub> with LogR<sub>s</sub>**

$N$	$A_{\text{FAERS}}$			$A_*$		
	$\overline{\text{rec}}_t$	$\overline{\text{prec}}_t$	$\overline{\text{acc}}_t$	$\overline{\text{rec}}_t$	$\overline{\text{prec}}_t$	$\overline{\text{acc}}_t$
5	<b>0.2358</b>	<b>0.2359</b>	0.2915	<b>0.2613</b>	<b>0.2615</b>	0.2821
	0.2324	0.2325	<b>0.2929</b>	0.2589	0.2592	<b>0.2832</b>
10	<b>0.2497</b>	<b>0.2497</b>	0.3076	<b>0.3469</b>	<b>0.3467</b>	0.3419
	0.2467	0.2467	<b>0.3110</b>	0.3434	0.3432	<b>0.3439</b>
20	<b>0.2885</b>	<b>0.2887</b>	0.2762	<b>0.4403</b>	<b>0.4407</b>	0.4016
	0.2694	0.2695	<b>0.3289</b>	0.4265	0.4273	<b>0.4135</b>

Column of “N” represents the number of recommended drugs. Columns of “ $A_{\text{FAERS}}$ ” and “ $A_*$ ” represent that “ $A_{\text{FAERS}}$ ” and “ $A_*$ ” are used as  $A_{\text{pool}}$ , respectively. The **bold** performance is the best under the corresponding metrics in each column.