

# Algorithm Pseudocode

## GPU-Accelerated Semantic Severity Recalibration

### 1 Main Recalibration Algorithm

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**Algorithm 1** Semantic Severity Recalibration (GPU-Accelerated)

Dataset  $\mathcal{D} = \{(desc^{(i)}, d_1^{(i)}, d_2^{(i)}, \hat{y}^{(i)}, c^{(i)})\}_{i=1}^N$  Severity prototypes  $\mathcal{P} = \{P_C, P_{Ma}, P_{Mo}, P_{Mi}\}$   
Sentence encoder  $f_\theta$  (all-MiniLM-L6-v2) Drug class table  $\mathcal{R}$  Weights  $w_s = 0.45$ ,  
 $w_c = 0.25$ ,  $w_d = 0.30$  Target distribution  $\mathcal{T} = [0.05, 0.25, 0.60, 0.10]$  Recalibrated  
predictions  $\{\tilde{y}^{(i)}\}_{i=1}^N$  **Phase 1:** Batch encode all descriptions on GPU  $\mathbf{E} \leftarrow GPUBatchEncode_{f_\theta}, \{desc^{(i)}\}_{i=1}^N$  Shape:  $N \times d$   $\mathbf{E} \leftarrow \mathbf{E}/\|\mathbf{E}\|$  Normalize embeddings  
**Phase 2:** Compute prototype centroids  $k \in \{C, Ma, Mo, Mi\}$   $\mathbf{c}_k \leftarrow \text{mean}(f_\theta(P_k))$   
Class centroid  $\mathbf{c}_k \leftarrow \mathbf{c}_k/\|\mathbf{c}_k\|$   $\mathbf{C} \leftarrow [\mathbf{c}_C; \mathbf{c}_{Ma}; \mathbf{c}_{Mo}; \mathbf{c}_{Mi}]$  Centroid matrix:  $4 \times d$  **Phase 3:** Vectorized similarity computation  $\mathbf{S}_{sem} \leftarrow \mathbf{E} \cdot \mathbf{C}^T$  Cosine similarity:  $N \times 4$   
 $\mathbf{s}_{sem} \leftarrow \text{ThresholdToScores}_{\mathbf{S}_{sem}}$  Map to numeric **Phase 4:** Parallel drug class  
scoring  $\mathbf{s}_{drug} \leftarrow \text{ParallelDrugClassRisk}\{(d_1^{(i)}, d_2^{(i)})\}, \mathcal{R}$  **Phase 5:** Confidence ad-  
justment  $\mathbf{s}_{conf} \leftarrow \text{AdjustConfidence}\{\hat{y}^{(i)}\}, \{c^{(i)}\}$  **Phase 6:** Weighted combination  
 $\mathbf{s}_{final} \leftarrow w_s \cdot \mathbf{s}_{sem} + w_c \cdot \mathbf{s}_{conf} + w_d \cdot \mathbf{s}_{drug}$  **Phase 7:** Quantile-based target calibration  
 $\{\tilde{y}^{(i)}\} \leftarrow \text{QuantileCalibrates}_{final}, \mathcal{T} \{\tilde{y}^{(i)}\}_{i=1}^N$

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## 2 Semantic Similarity Scoring

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**Algorithm 2** ThresholdToScores: Convert Similarities to Severity Scores

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Similarity matrix  $\mathbf{S} \in R^{N \times 4}$  (columns: Contra, Major, Mod, Minor) Thresholds  $\tau_C = 0.65$ ,  $\tau_{Ma} = 0.55$ ,  $\tau_{Mo} = 0.45$  Severity scores  $\mathbf{s} \in [1.5, 4.0]^N$   $\mathbf{s} \leftarrow \mathbf{1.5}$  Initialize all to Minor score  $i \in \{1, \dots, N\}$  Vectorized in practice  $S_{i,0} \geq \tau_C$   $s_i \leftarrow 4.0$  Contraindicated  $S_{i,1} \geq \tau_{Ma}$   $s_i \leftarrow 3.2$  Major  $S_{i,2} \geq \tau_{Mo}$   $s_i \leftarrow 2.0$  Moderate  $\mathbf{s}$

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## 3 Quantile-Based Target Calibration

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**Algorithm 3** QuantileCalibrate: Exact Target Distribution Matching

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Final scores  $\mathbf{s}_{final} \in R^N$  Target distribution  $\mathcal{T} = [t_C, t_{Ma}, t_{Mo}, t_{Mi}]$  Severity labels  $\{\tilde{y}^{(i)}\}_{i=1}^N$   $\pi \leftarrow ArgSort\mathbf{s}_{final}$ , descending Sort indices by score  $N \leftarrow |\mathbf{s}_{final}|$  Compute category boundaries from targets  $n_C \leftarrow \lfloor N \cdot t_C \rfloor$  Top 5%  $n_{Ma} \leftarrow \lfloor N \cdot t_{Ma} \rfloor$  Next 25%  $n_{Mi} \leftarrow \lfloor N \cdot t_{Mi} \rfloor$  Bottom 10%  $n_{Mo} \leftarrow N - n_C - n_{Ma} - n_{Mi}$  Remaining 60% Assign severities by rank  $\tilde{\mathbf{y}} \leftarrow \mathbf{0}^N$   $\tilde{y}_{\pi[1:n_C]} \leftarrow$  “Contraindicated”  $\tilde{y}_{\pi[n_C+1:n_C+n_{Ma}]} \leftarrow$  “Major”  $\tilde{y}_{\pi[n_C+n_{Ma}+1:N-n_{Mi}]} \leftarrow$  “Moderate”  $\tilde{y}_{\pi[N-n_{Mi}+1:N]} \leftarrow$  “Minor”  $\tilde{\mathbf{y}}$

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## 4 Confidence Adjustment

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**Algorithm 4** AdjustConfidence: Penalize Low-Confidence High-Severity

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Original predictions  $\{\hat{y}^{(i)}\}$  and confidences  $\{c^{(i)}\}$  Thresholds  $\tau_C = 0.65$ ,  $\tau_{Ma} = 0.50$  Confidence-adjusted scores  $\mathbf{s}_{conf}$   $\phi \leftarrow \{C : 4, Ma : 3, Mo : 2, Mi : 1\}$  Label to score map  $i \in \{1, \dots, N\}$   $s_i \leftarrow \phi(\hat{y}^{(i)})$  Base score from prediction  $\hat{y}^{(i)}$  = “Contraindicated” **and**  $c^{(i)} < \tau_C$   $s_i \leftarrow 3.0$  Downgrade uncertain contraindicated  $\hat{y}^{(i)} \in \{\text{“Contra”}, \text{“Major”}\}$  **and**  $c^{(i)} < \tau_{Ma}$   $s_i \leftarrow 2.5$  Partial downgrade  $\mathbf{s}_{conf}$

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## 5 Drug Class Risk Assessment

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**Algorithm 5** ParallelDrugClassRisk: Pharmacological Risk Scoring

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Drug pairs  $\{(d_1^{(i)}, d_2^{(i)})\}_{i=1}^N$  High-risk classes  $\mathcal{R} = \{\text{anticoag}, \text{antiplate}, \text{QT}, \text{MAOI}, \dots\}$  Drug class scores  $\mathbf{s}_{drug}$  Parallel execution on 24 CPU cores **parallel for**  $i \in \{1, \dots, N\}$  **do**  $C_1 \leftarrow GetClassesd_1^{(i)}, \mathcal{R}$   $C_2 \leftarrow GetClassesd_2^{(i)}, \mathcal{R}$   $O \leftarrow C_1 \cap C_2$  Overlap “MAOI”  $\in C_1 \wedge$  “Serotonergic”  $\in C_2$   $s_i \leftarrow 4.0$  Fatal combination “MAOI”  $\in O$   $s_i \leftarrow 4.0$  “anticoag”  $\in O \vee$  “QT”  $\in O$   $s_i \leftarrow 3.5$   $C_1 \neq \emptyset \wedge C_2 \neq \emptyset$   $s_i \leftarrow 3.0$   $C_1 \neq \emptyset \vee C_2 \neq \emptyset$   $s_i \leftarrow 2.5$   $s_i \leftarrow 2.0$  **end parallel for**  $\mathbf{s}_{drug}$

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## 6 Complexity Analysis

Phase	Time Complexity	Actual Time
GPU Batch Encoding	$O(N \cdot d/B)$	45.5s
Centroid Computation	$O( \mathcal{P}  \cdot d)$	<1s
Similarity Matrix	$O(N \cdot 4)$	0.04s
Drug Class Scoring	$O(N/P)$	0.7s
Confidence Adjustment	$O(N)$	<1s
Quantile Calibration	$O(N \log N)$	<1s
<b>Total</b>	$O(N \cdot d/B + N \log N)$	<b>49.2s</b>

Table 1: Complexity analysis for  $N = 759,774$ ,  $d = 384$ ,  $B = 8192$ ,  $P = 24$  cores

## 7 Performance Summary

- **GPU:** NVIDIA RTX PRO 5000 (48GB VRAM)
- **CPU:** 24 cores for parallel drug class scoring
- **Memory:** 124GB RAM
- **Throughput:** 15,454 interactions/second
- **Embedding Rate:** 16,696 descriptions/second
- **Total Time:** 49.2 seconds for 759,774 interactions
- **Target Match:** Exact (0% deviation from literature targets)