

# Automatically Labeling Clinical Trial Outcomes: A Large-Scale Benchmark for Drug Development

Chufan Gao\*, Jathurshan Pradeepkumar\*, Trisha Das\*, Shivashankar Thati, and Jimeng Sun



Project Page



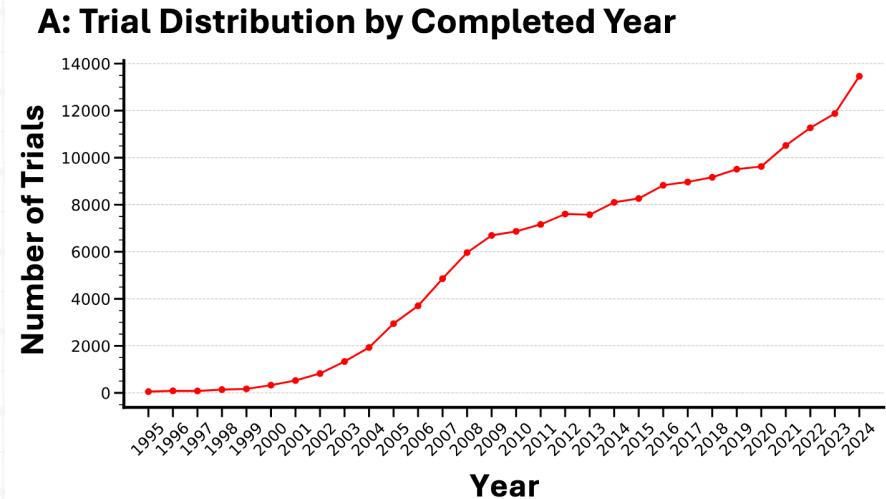
Paper



Code

# Background

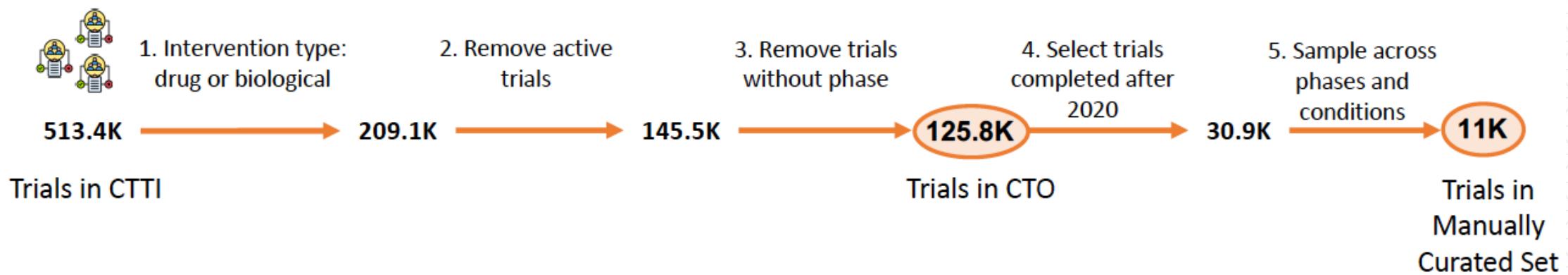
- Drug discovery and development are expensive, with clinical trial results vital for regulatory approval and patient care.
- Large-scale, high-quality clinical trial outcome data remains limited.
  - Hindering the development of predictive models
- Dynamic & Rapidly Growing Data
  - Clinical trial data grows rapidly and is affected by diverse external factors (e.g., COVID-19, regulatory changes).
  - Frequent label updates are needed, but **manual labeling is impractical at scale**.



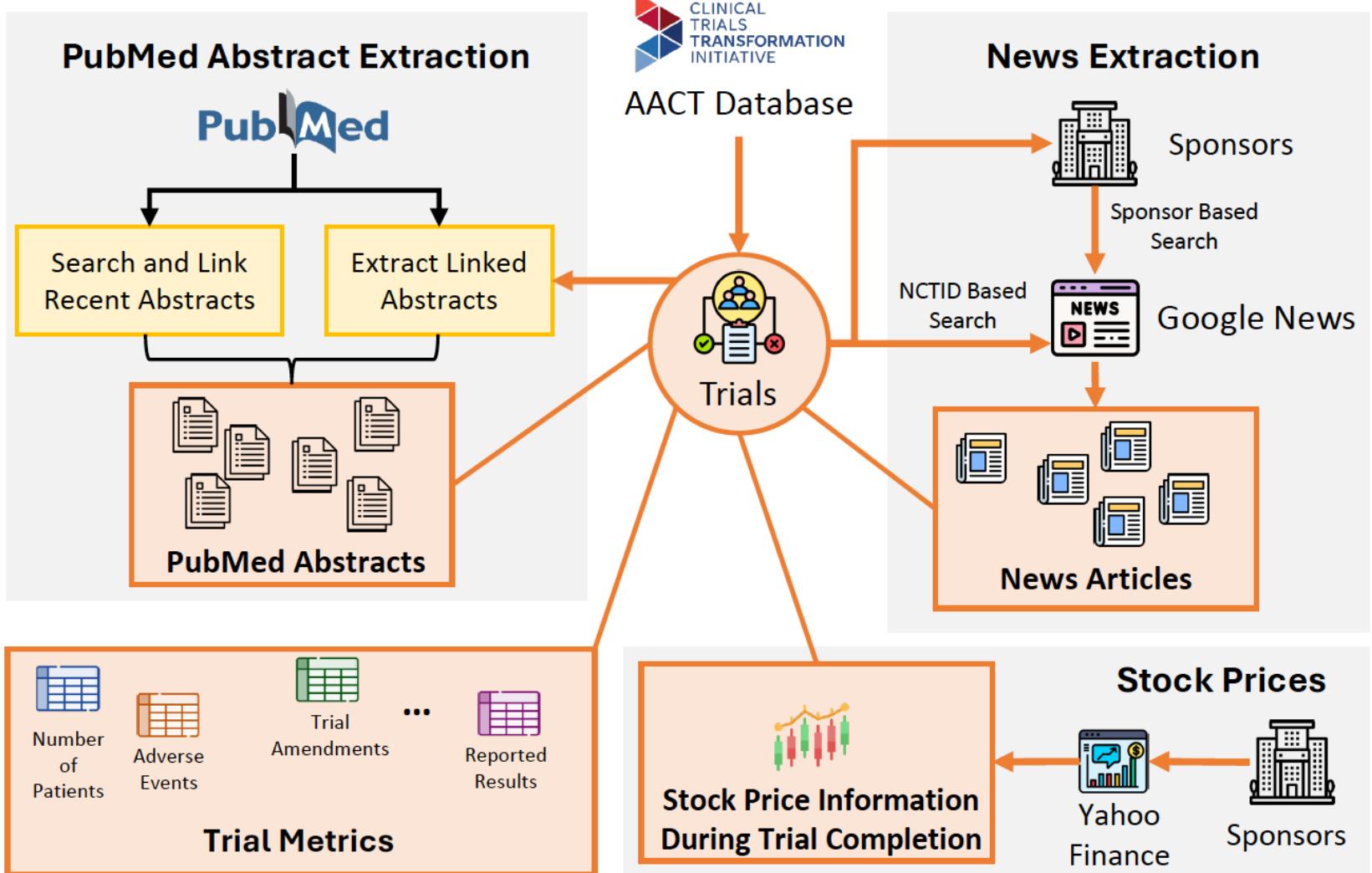
# Clinical Trial Outcome Benchmark (CTO)

- Clinical Trial Outcome (CTO) benchmark, a fully reproducible, regularly updated, large-scale repository encompassing approximately ~125K drug and biologics trials.

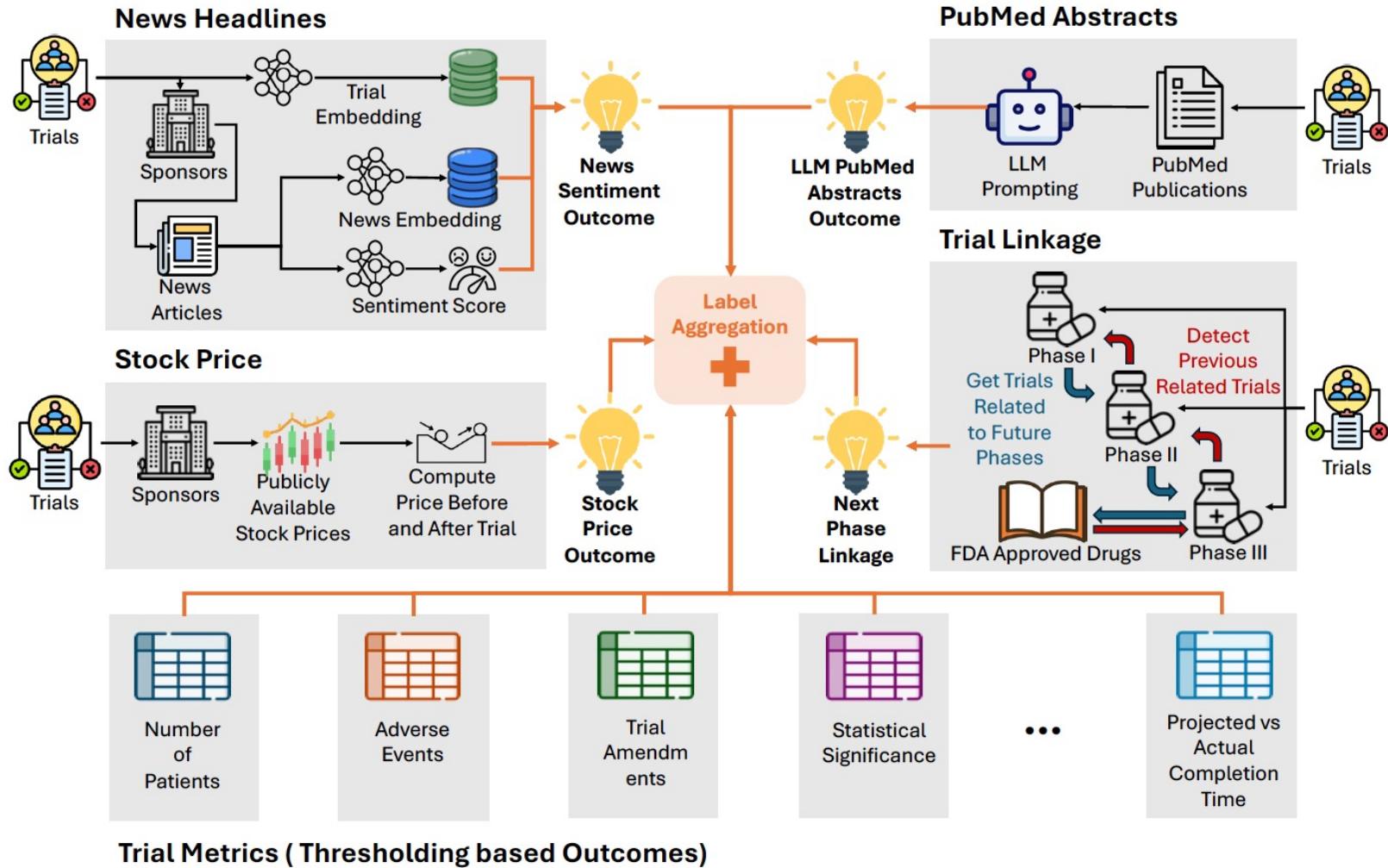
- A comprehensive trial knowledge base
- Automated labeling framework based on aggregation of weak labeling function
- Manually curated around 11K trials.



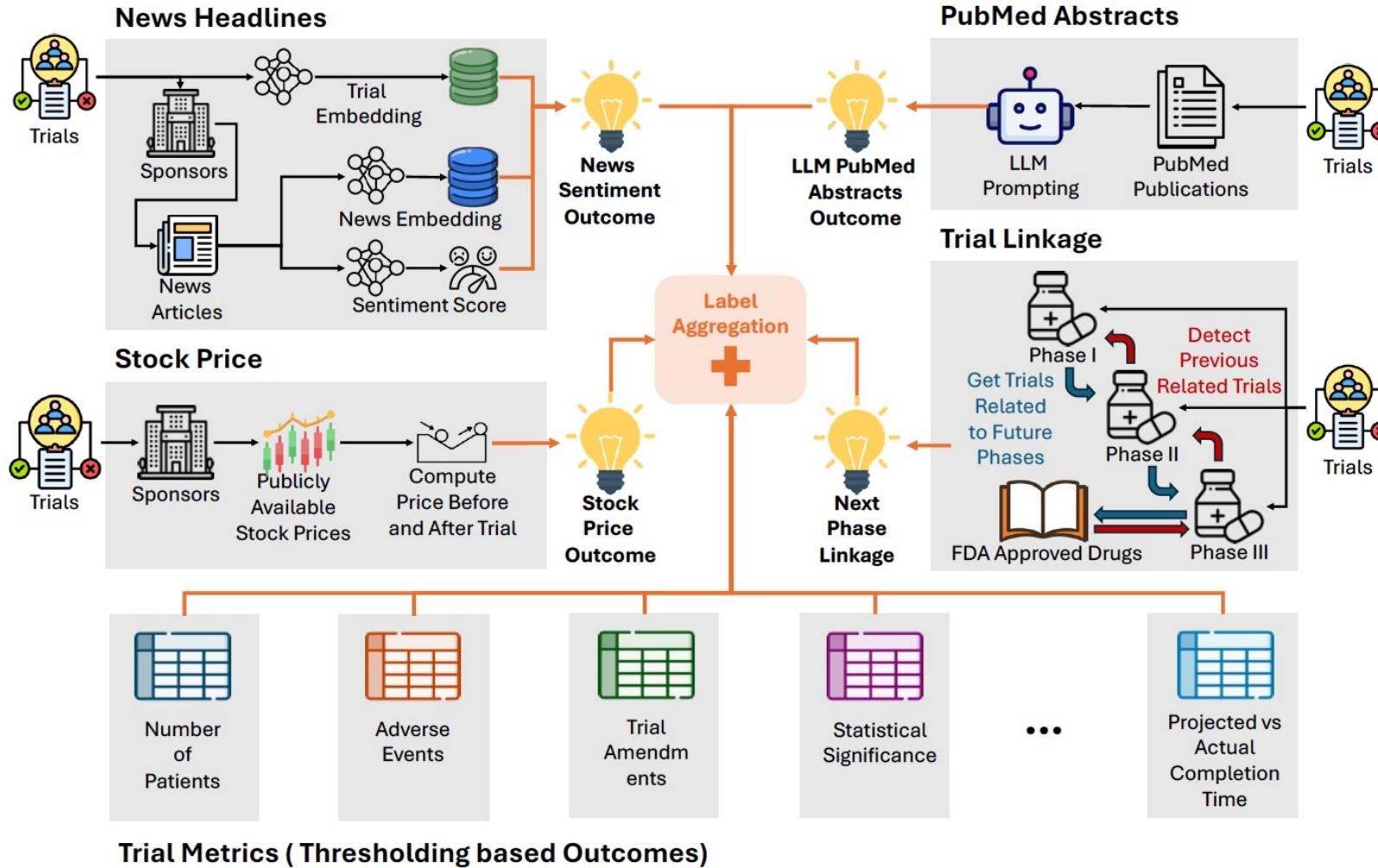
# Trial Knowledge Base



# CTO Automated Labeling Framework



# CTO Automated Labeling Framework



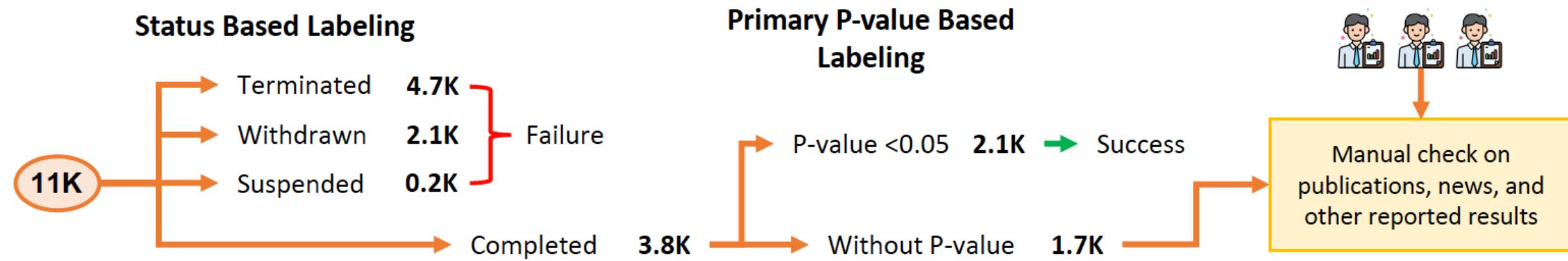
Phase	Aggregation Method	F1	$\kappa$
I	$CTO_{MV}$	0.726	0.490
	$CTO_{DP}$	0.870	0.700
	$CTO_{RF}$	<b>0.913</b>	<b>0.790</b>
II	$CTO_{MV}$	0.689	0.430
	$CTO_{DP}$	0.856	0.623
	$CTO_{RF}$	<b>0.878</b>	<b>0.693</b>
III	$CTO_{MV}$	0.904	0.606
	$CTO_{DP}$	0.921	0.582
	$CTO_{RF}$	<b>0.941</b>	<b>0.710</b>
All	$CTO_{MV}$	0.793	0.529
	$CTO_{DP}$	0.884	0.646
	$CTO_{RF}$	<b>0.909</b>	<b>0.729</b>

Agreement of CTO automated labels with human-labeled TOP[1] dataset

## Trial Metrics ( Thresholding based Outcomes)

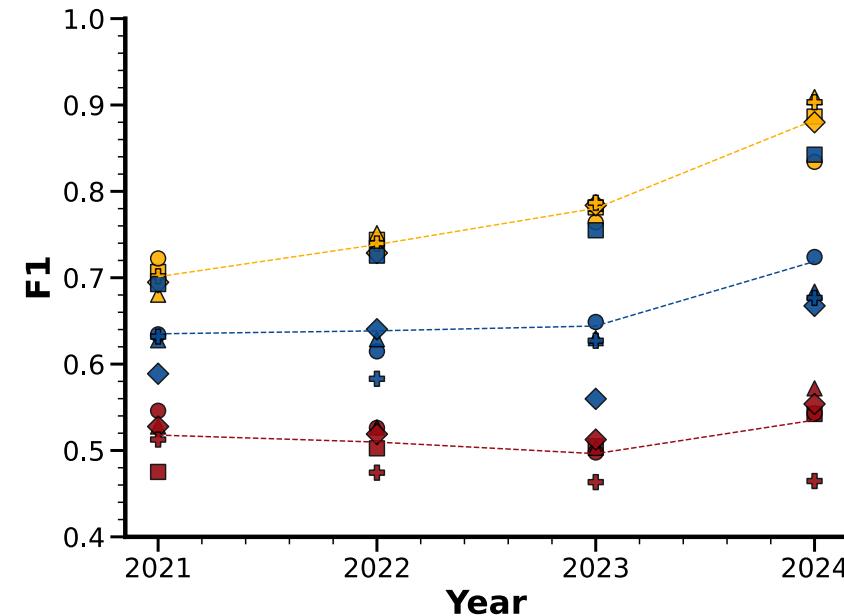
[1] Fu, Tianfan, et al. "Hint: Hierarchical interaction network for clinical-trial-outcome predictions." *Patterns* 3.4 (2022).

# Manual Curation Process

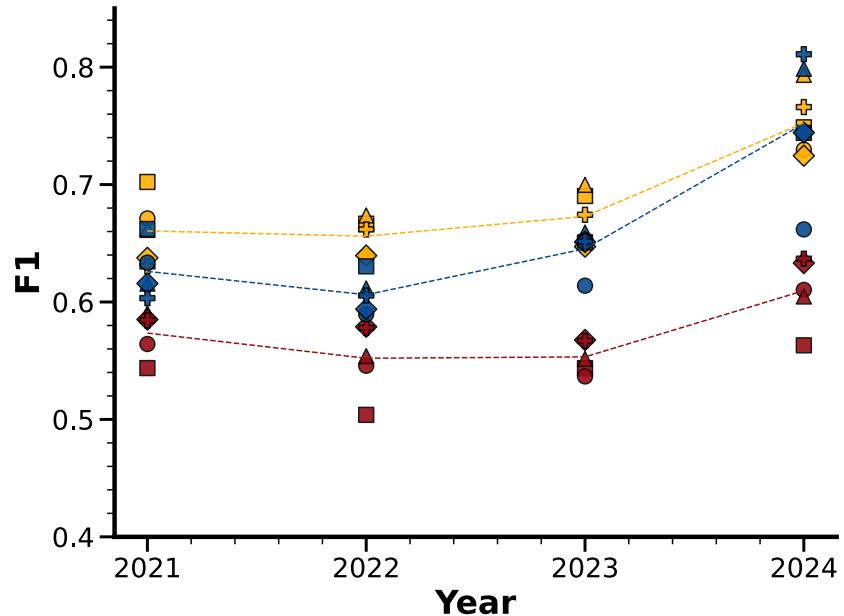


# Which labels are better?

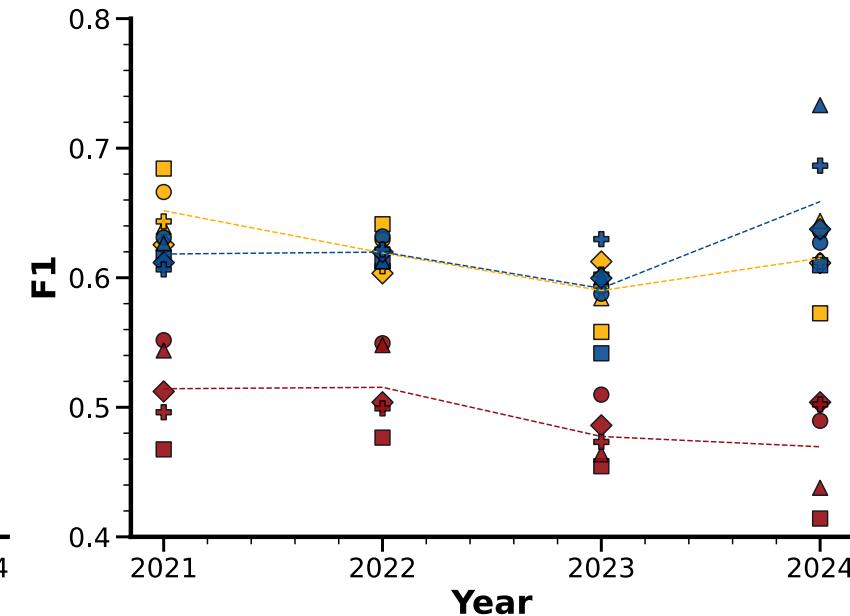
Phase 1



Phase 2



Phase 3



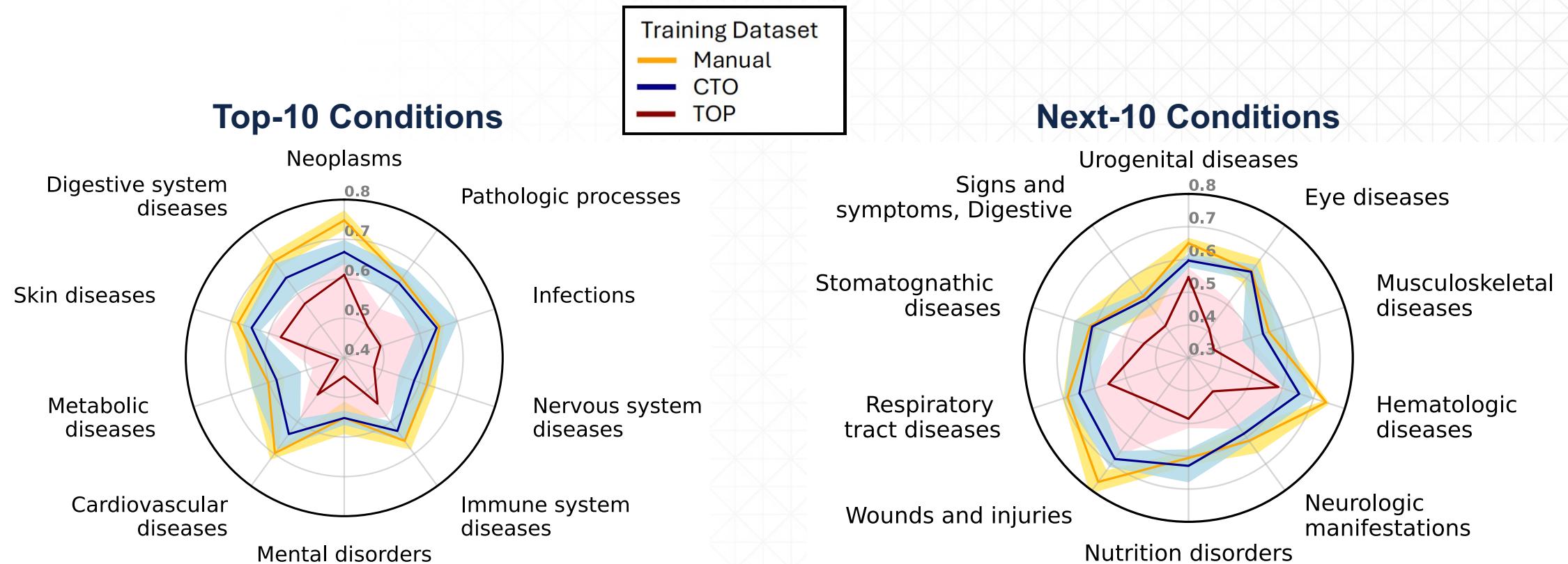
- Manual : Updated human - annotated data  
CTO : From our automated labeling framework  
TOP : Past benchmark on clinical trial outcomes [1]

[1] Fu, Tianfan, et al. "Hint: Hierarchical interaction network for clinical-trial-outcome predictions." *Patterns* 3.4 (2022).

Training Dataset - Baseline Model

■ TOP - Random Forest	■ Manual - Random Forest	■ CTO - Random Forest
● TOP - Logistic Regression	● Manual - Logistic Regression	● CTO - Logistic Regression
▲ TOP - XGBoost	▲ Manual - XGBoost	▲ CTO - XGBoost
◆ TOP - MLP	◆ Manual - MLP	◆ CTO - MLP
✚ TOP - SVM	✚ Manual - SVM	✚ CTO - SVM
--- TOP (Mean)	--- Manual (Mean)	--- CTO (Mean)

# Which labels are better?



Manual	: Updated human - annotated data
CTO	: From our automated labeling framework
TOP	: Past benchmark on clinical trial outcomes [1]

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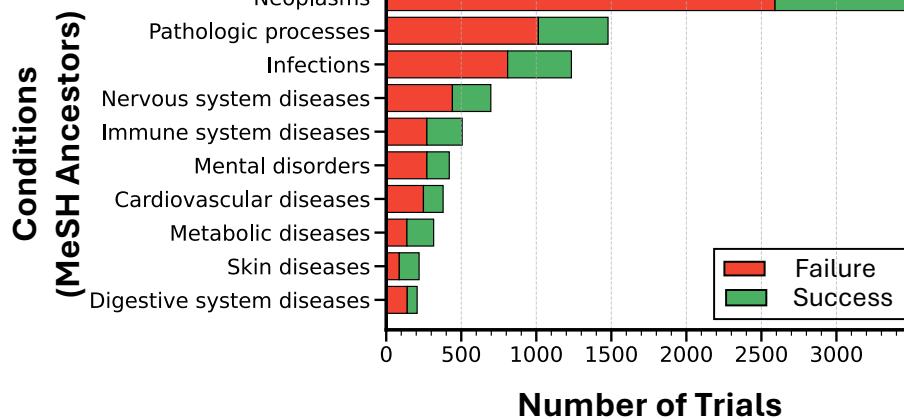
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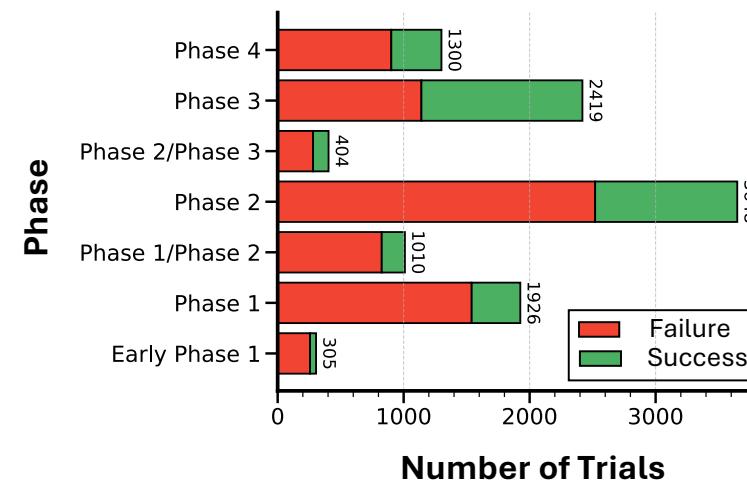
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# Manual Curated Trial Outcome Benchmark

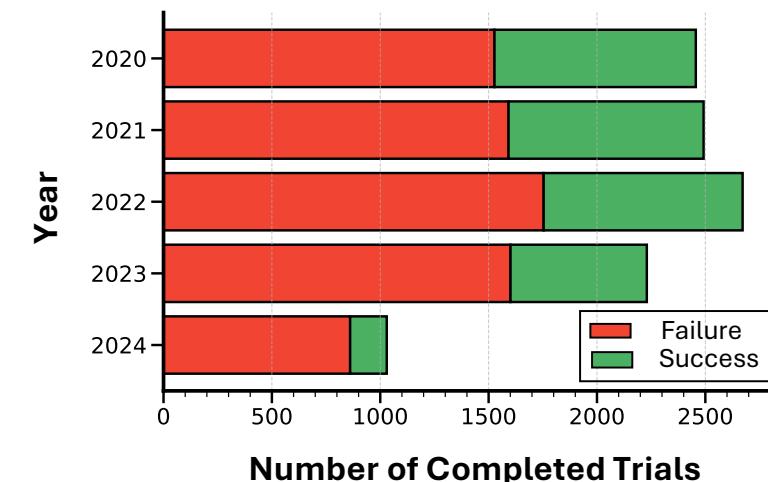
A: Top 10 Disease Distribution



B: Phase Distribution

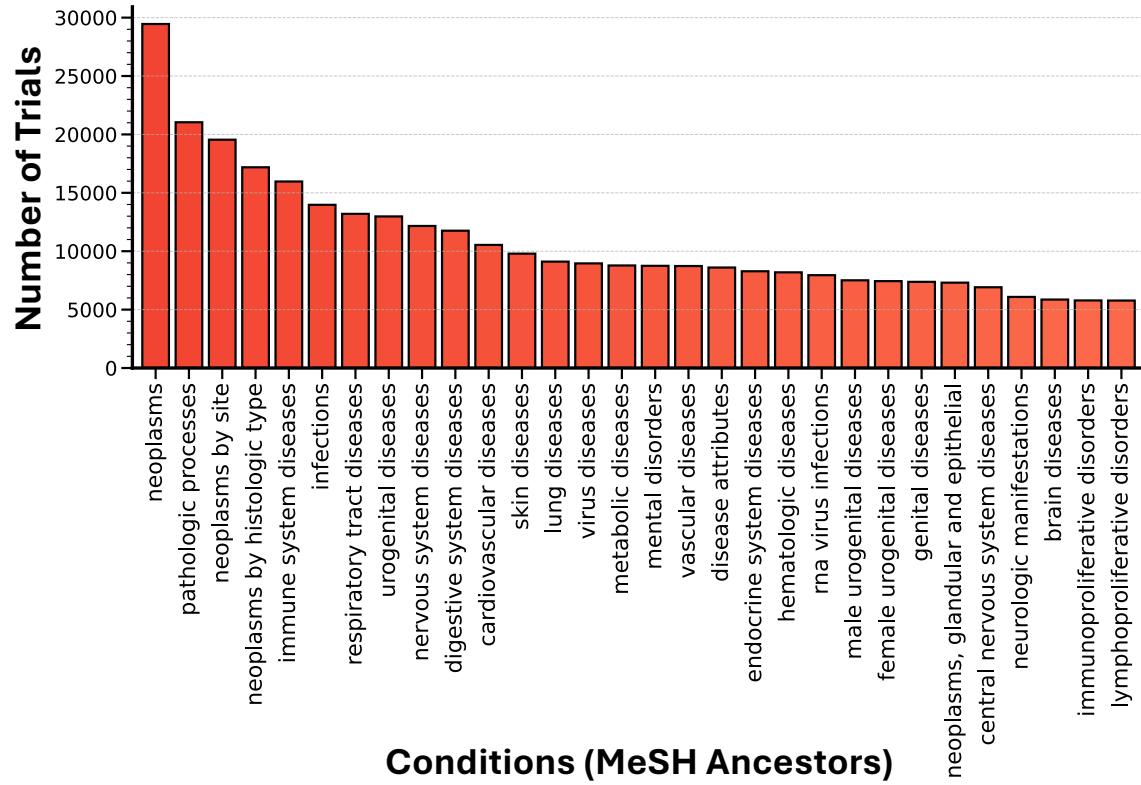


C: Distribution by Completed Year

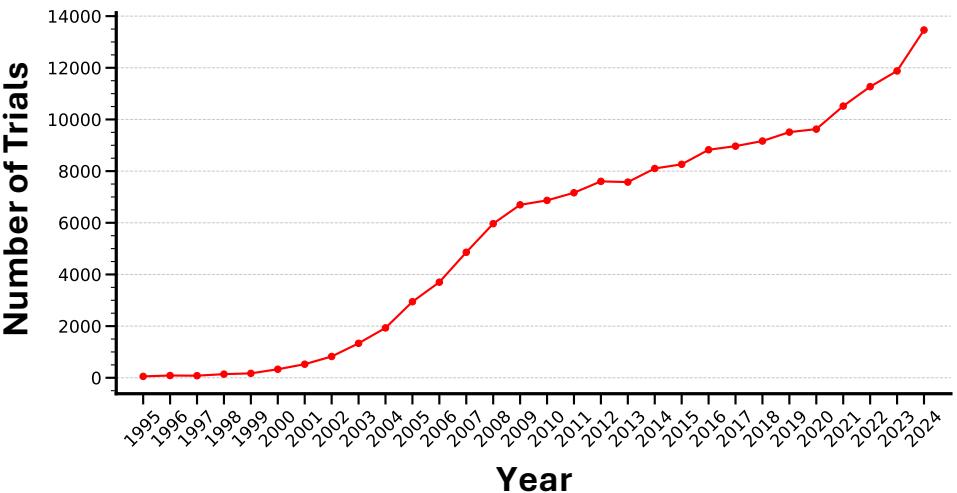


# CTO Statistics

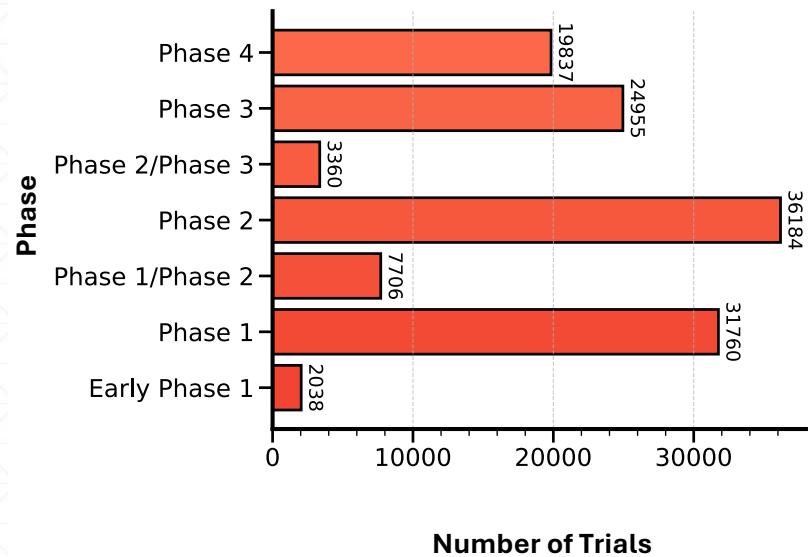
B: Frequency of Top 30



A: Trial Distribution by Completed Year

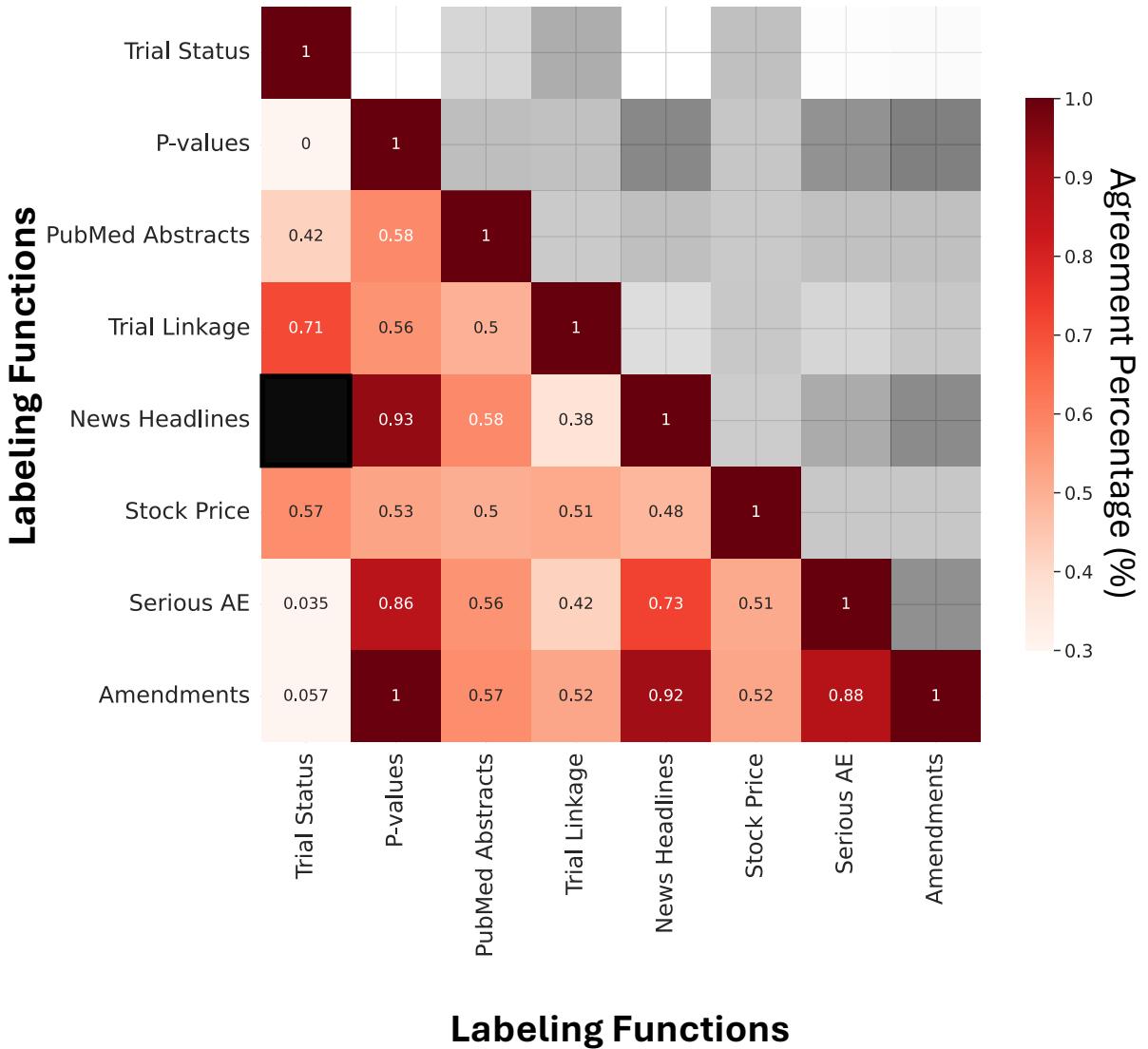


D: Phase Distribution



# Agreement between Labeling Functions in CTO

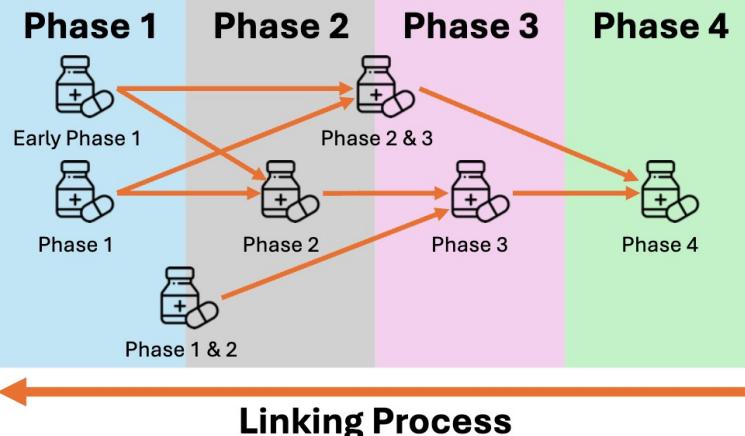
C: Agreement between Labeling Functions in CTO



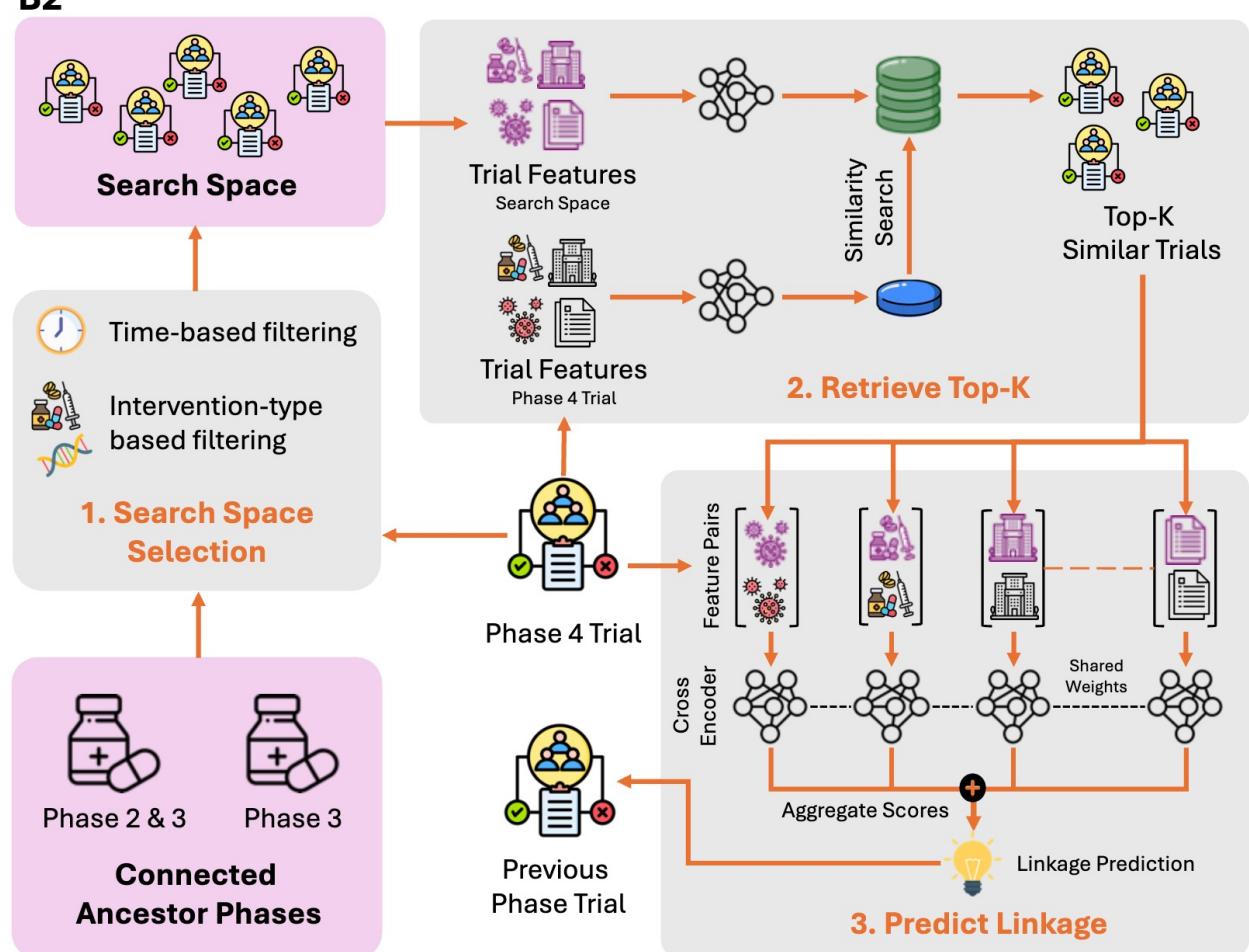
# Additional Method / Weak Labeling Slides

## B. Trial Linkage Algorithm

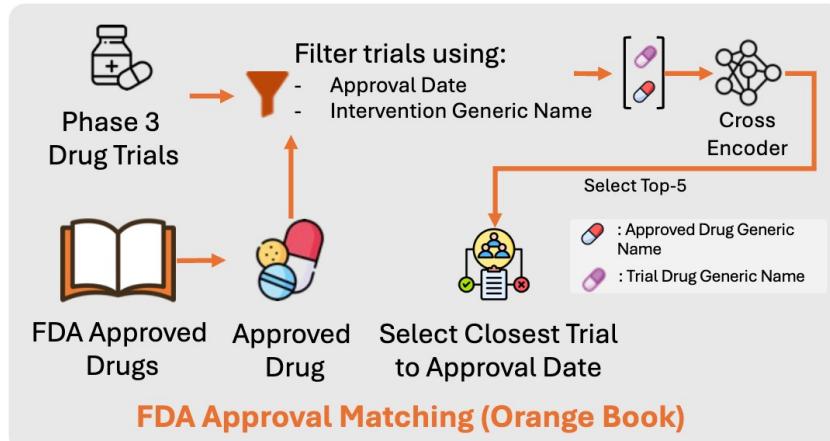
B1



B2



B3



# Additional Method / Weak Labeling Slides

