# **Table-to-text Generation for Biomedical Causal Inference**

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# FDA Adverse Event Reporting System (FAERS)

- Over 2 million reports every year
- Example Report:

CaseID	Gender	Age	PSD	ADE	•••	
20222515	Female	71	Nicotine, Acetaminophen	Anemia, Headache	•••	

PSD = Primary Suspect Drugs

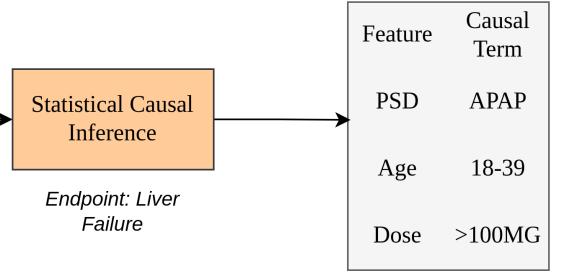
ADE = Adverse Drug Effects

## **Conventional CI pipeline**

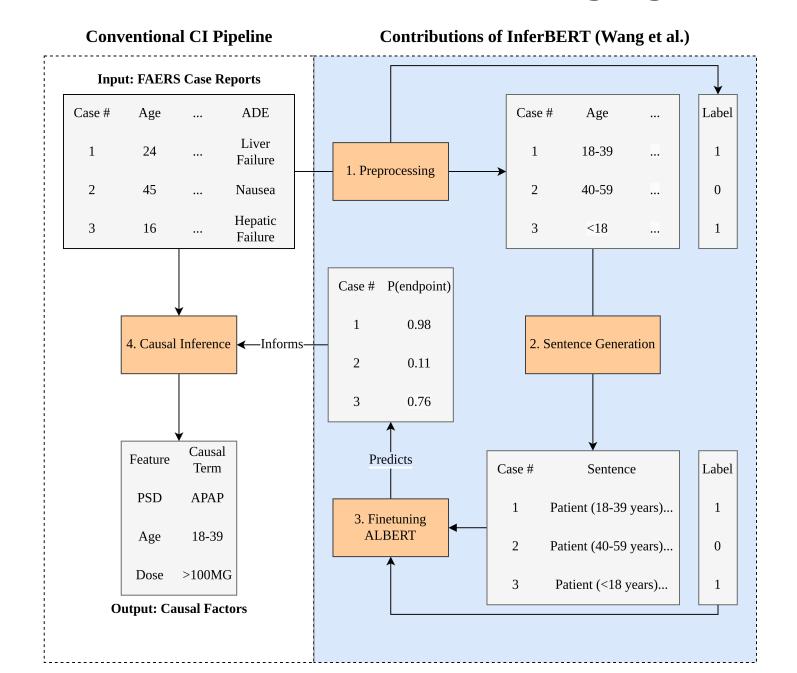
### **Input: Pharmacovigilance Case Reports**

#### Case # **ADE** Age • • • Liver 1 24 ••• Failure 2 45 Nausea • • • Hepatic 16 3 ••• Injury

### **Output: Causal Factors**



### InferBERT informs causal inference with a language model (ALBERT)

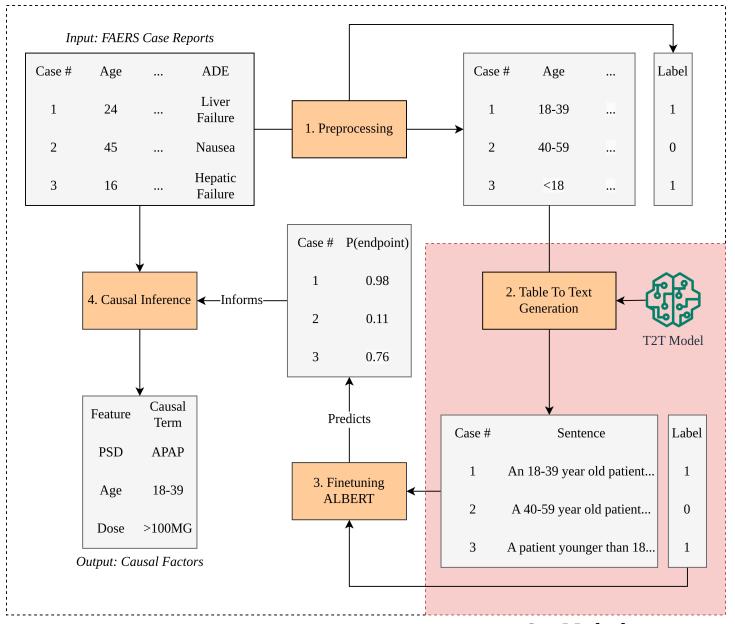


# Analgesics-Induced acute liver failure 114.3 Acetaminophen | 20.4 12.4 Death 18-39 6.1 Female

### **InferBERT Results**

Feature	Term	Z- score	
PSD	APAP	114.95	
Outcome	Death	77.94	
Age	18-39	27.29	
Indication	Suicide Attempt	16.85	
Gender	Female	10.26	
Dose	>100mg	5.01	

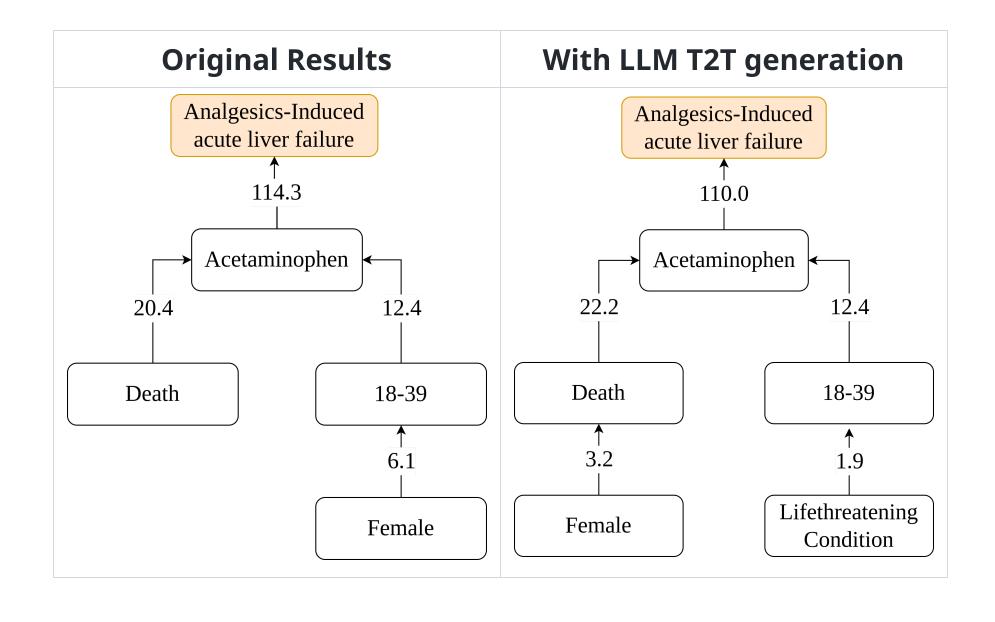
### We modify InferBERT using Table-to-text generation

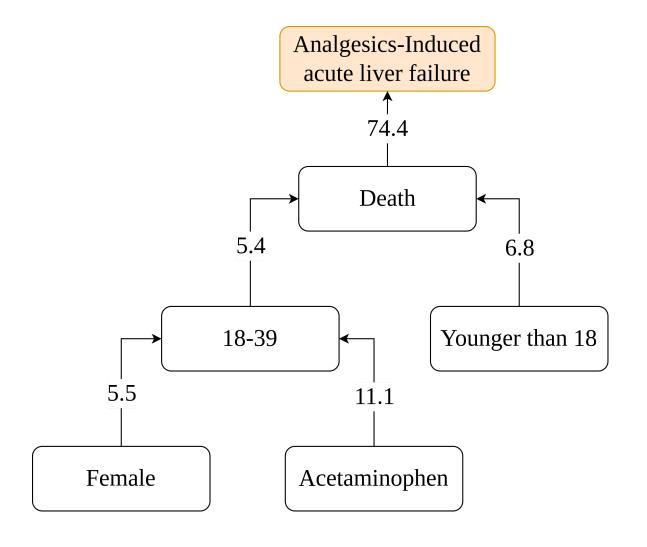


**Our Method** 

## Advantages of our method

- T2T generation can be applied to any input features This allows us to easily apply the framework to new studies or features
- Using a pre-trained language model, T2T is able to infer implicit relations between report terms
- These relations are made explicit in the generated sentences, improving endpoint prediction of the ALBERT model





### **With More Features**

Feature	Value	Z- score	
SSD	Diphen- hydramine	5.13	
Weight	50-70kg	5.75	
Route	Intravenous	15.55	

SSD= Secondary Suspect Drug

### **Future Work**

- Applying this framework to further case studies
- What are the limits for the number of features/terms we can study in one go?
- What would be the impact of using prompting instead of finetuning for endpoint prediction?
  - $\circ$  Can we use a single model from report  $\rightarrow$  prediction?