



DD.ai

A Platform for Drug-Drug Interaction Prediction



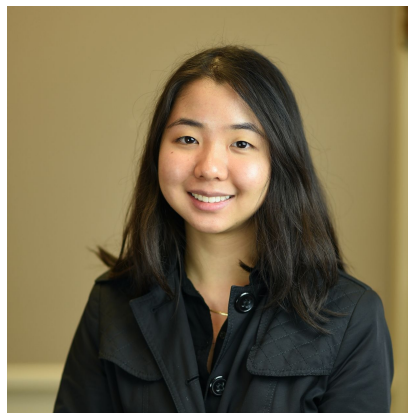


Our Team



Mai La

Data Scientist
Data Engineer | MLOps



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Product Manager | UI



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Data Scientist
ML Engineer | UI



01

Impact

Opportunity
Target Users

02

Product

Quick Demo

03

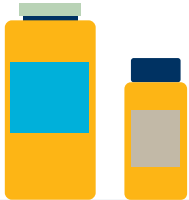
Approach

Details on data
Scoping
ML set up

04

Results

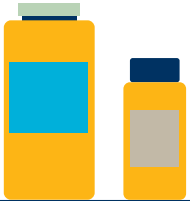
Evaluation
Future Work
Conclusion





01

Impact





Drug - Drug interactions (DDI) can have severe effects



52% of adverse drug effects are **preventable**¹



Account for **18.3%** of all adverse drug effects²



22.4% of adults in the US use **5+** medications³

¹ Source: [Hakkarainen et al., 2012](#)

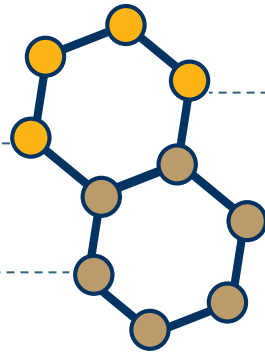
² Source: [Jiang et al., 2022](#)

³ Source: [Hales et al., 2019](#)

Drug Interaction Testing is Expensive

\$60 Billion
2023 Global Spending⁴

60% increase
Drug approvals
2010-2019⁶



\$133 Billion
2032 Global Spending⁵

4-5 Years
For testing full panel and
reporting⁷

⁴ Source: [Precedence Research](#)

⁶ Source: [CBO](#)

⁵ Source: [Precedence Research](#)

⁷ Source: User Interviews

User Interviews



Pharmacist - Private

- Manually find DDI



Supplement Co CEO

- Report for FDA
- Lots of testing



Org Chem PhD

- Chemical Formula
- Mechanism of Action



Pharmacist - CVS

- Have built in database for searching





DD.ai



Optimize DDI testing and assist drug companies to bring new drugs to market



Input:

- SMILES (Simplified Molecular-Input Line-Entry System) Drug structure

```
COC1=CC2=C(C=C1)N=C(N2)S(=O)CC1=NC=C(C(C)OC)=C1C
```

- Mechanism of Action Pathway

This drug stops gastric acid secretion by selective inhibition of the H⁺/K⁺ ATPase enzyme system...



Output:

- Prediction Probability of Positive Interactions
- Breakdown of supplement vs medication
- Downloadable table with drug list and probabilities



02 Product



Demo

Product Feedback



Dr. Han Dang, PharmD
Pharmacy Manager

Impressive that AI could extract DDI for new drug development.



Mai Do, RPh., MBA
Pharmacy Manager, CVS

I was amazed by how fast this tool can help define possible interactions between a new drug with hundreds of available drugs at a very high level of accuracy. This definitely help drug discovery much faster and more efficient with a lower cost. Traditionally, the process of evaluating drug interactions might take months or years of experiments in vitro or in vivo. With DD.ai, it's less than 1 minute. Simply amazing!!!



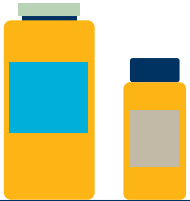
Dr. Tuan La, PharmD
CEO, Global Express Pharmacy

Good start for interaction checker. It would bring greater values to the users by serving a high accurate AI model which focus on a certain drug family like antibiotics or cancer treatment drugs.

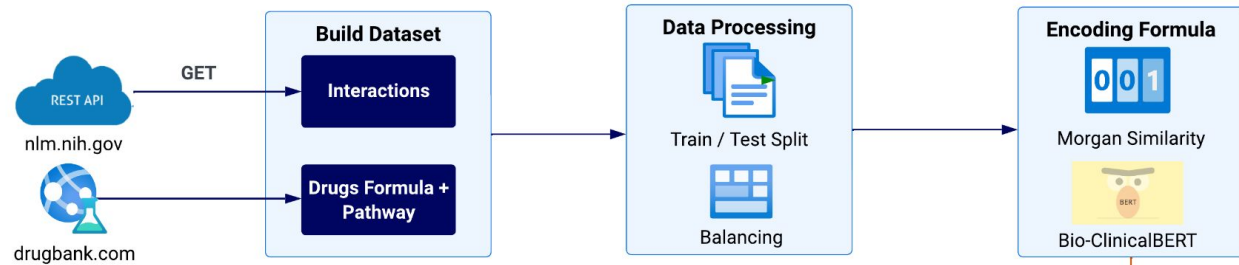


03

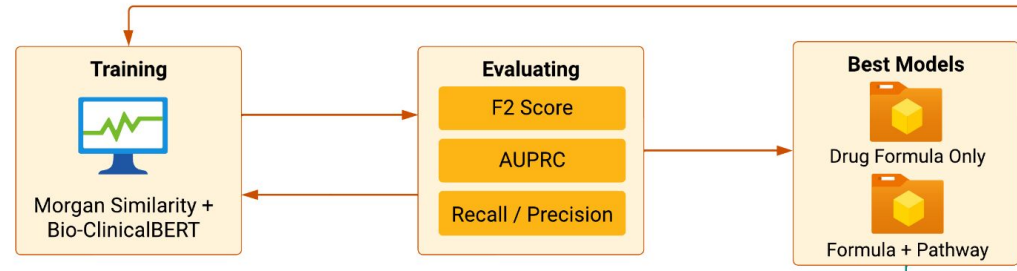
Approach



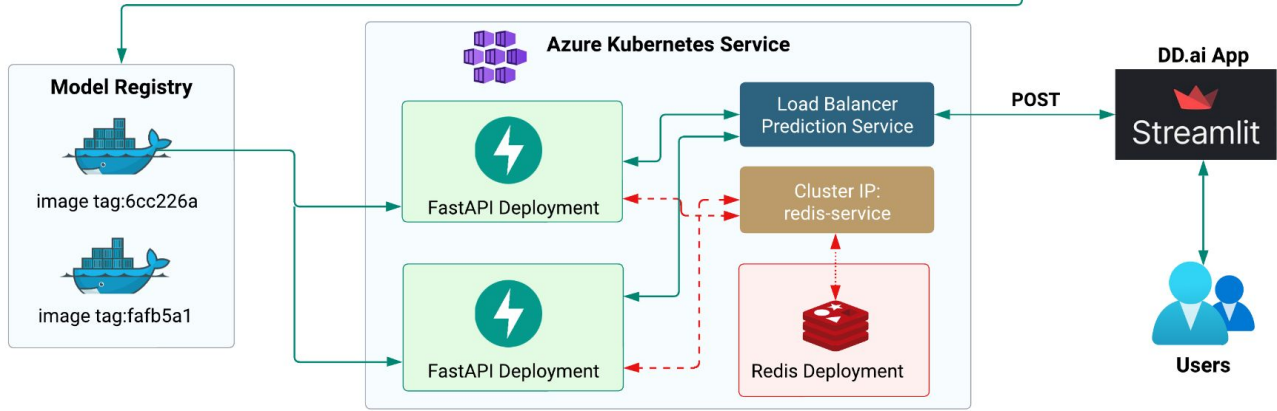
1 - Process Data



2 - Develop Model



3 - Production



Data Sources

DrugBank (Features):

- Drug SMILES
- Action Pathway

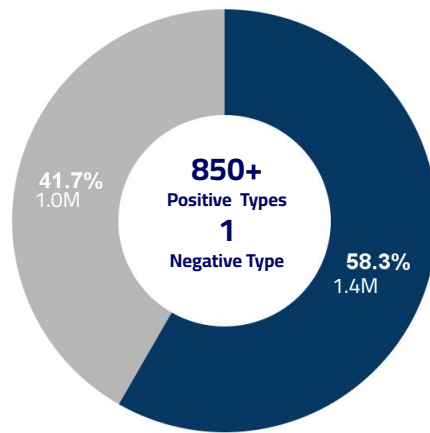
Median Word Count / Token Length



2,979 Drugs

NIH (Outcome):

- Drug Interactions



● Positive Interaction ● Negative Control



Key Considerations



850+
Interaction Types

Need to reduce
scope



SMILES
Chemical structures

Non-traditional
text encodings



512
Tokens

Limitation for NLP
BERT encodings



Drugbank
data

Limited to pairs of
drugs

Scope and Encodings



Scope

**850+ Class Prediction
Unfeasible**



Binary Classification

Text-based features

Morgan Similarity

Encode SMILES chemical structures into vectors

Google Pegasus PubMed

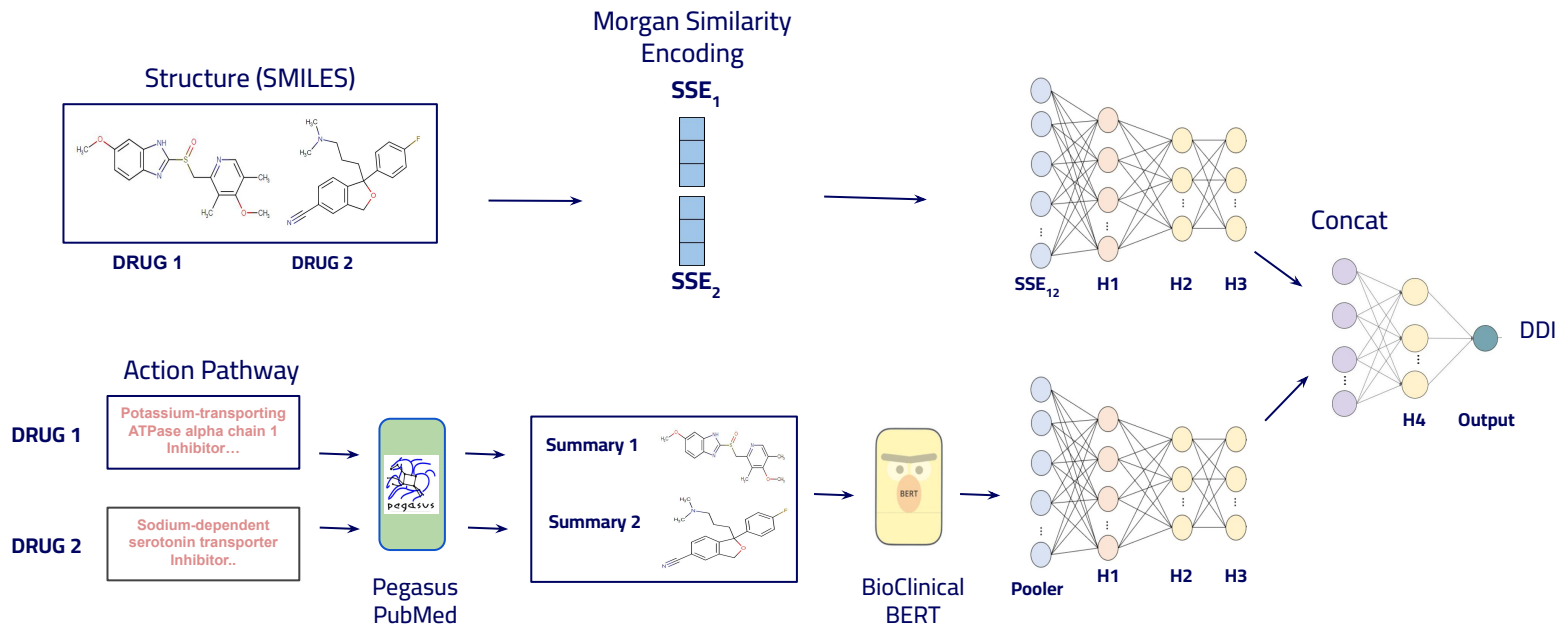
Summarize large text blocks of drug information

BioClinical BERT

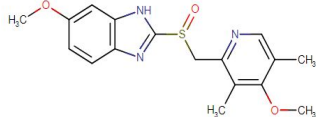
Encode above features for enhanced differentiability.



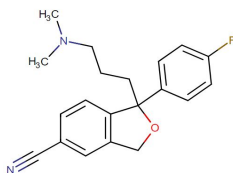
Architecture



DrugA = Omeprazole



DrugB = Citalopram



SMILES (Simplified Molecular-Input Line-Entry System)

COC1=CC2=C(C=C1)N=C(N2)S(=O)C1=NC=C(C)C(OC)=C1C

CN(C)CCCC1(OCC2=C1C=CC(=C2)C#N)C1=CC=C(F)C=C1

Morgan Fingerprints Encoding
(1024 bits)

00000100000001000010010000
000000010000100010000100...

00000000010000000100000010
000000000001000010000100...

Structure Similarity (SS)

Drug - Basis	Cosine Similarity
SS_{AC1}	0.161

Structure Similarity Encoding
(SSE) - 300 centroids from KMeans

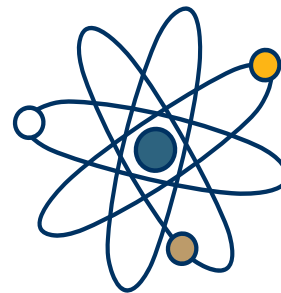
[0.161, 0.226, 0.204, 0.264, ...]

$SSE_A = [SS_{AC1}, SS_{AC2}, SS_{AC3}, \dots]$

[0.032, 0.196, 0.206, 0.160, ...]

$SSE_B = [SS_{BC1}, SS_{BC2}, SS_{BC3}, \dots]$

Morgan Similarity



Key Takeaways:

- Encode SMILES with Morgan Similarity
- Create 300 clusters of encodings
- Take cosine similarity of new SMILES with 300 clusters to reduce dimensionality

Text Encodings

Google Pegasus PubMed

Summarize large text blocks of drug information

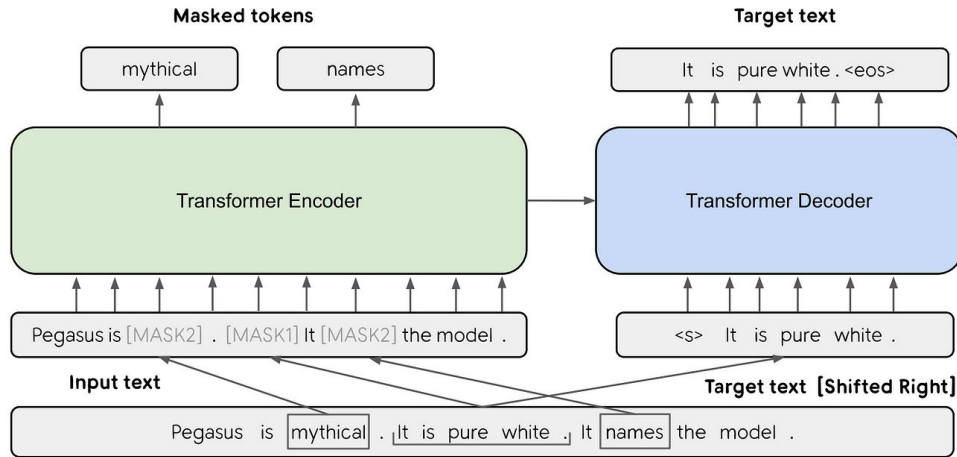


Image Source: <https://www.weak-learner.com/blog/2020/06/13/pegasus/>

BioClinical BERT

Encode features using BERT model pre-trained on biomedical texts

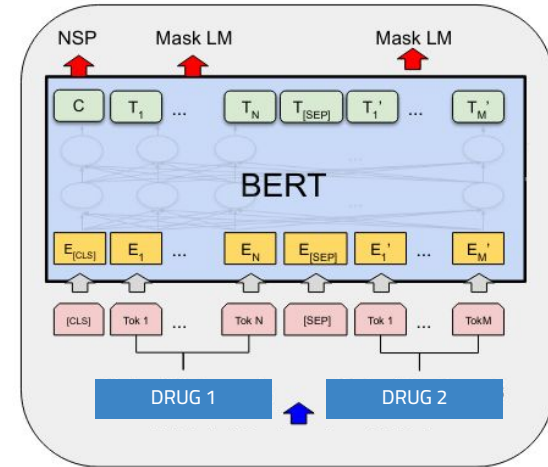
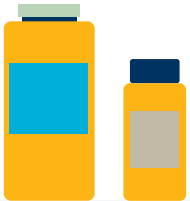


Image Source: <https://arxiv.org/pdf/1810.04805.pdf>



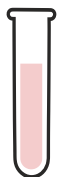
04

Results



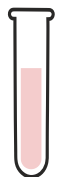
Model Registry

Baselines



Logistic Regression

F2: 70% AUPRC: 72%



XGBoost

F2: 70% AUPRC: 74%

Improvement



**Morgan Similarity
Neural Net**

F2: 73% AUPRC: 83%



**BioClinicalBERT
SMILES Only**

F2: 81% AUPRC: 86%

Final



**Morgan Similarity +
BioClinicalBERT
SMILES Only**

F2: 83% AUPRC: 81%

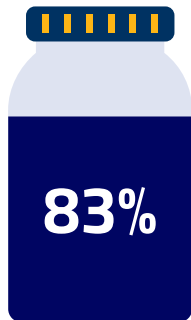


**Morgan Similarity +
BioClinicalBERT
SMILES & Pathway**

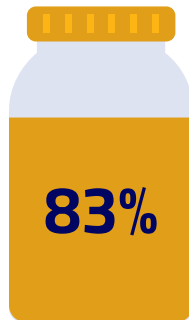
F2: 83% AUPRC: 83%

Evaluation

Morgan Similarity + BioClinicalBERT on SMILES & Pathway



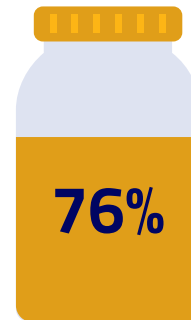
F2



AUPRC

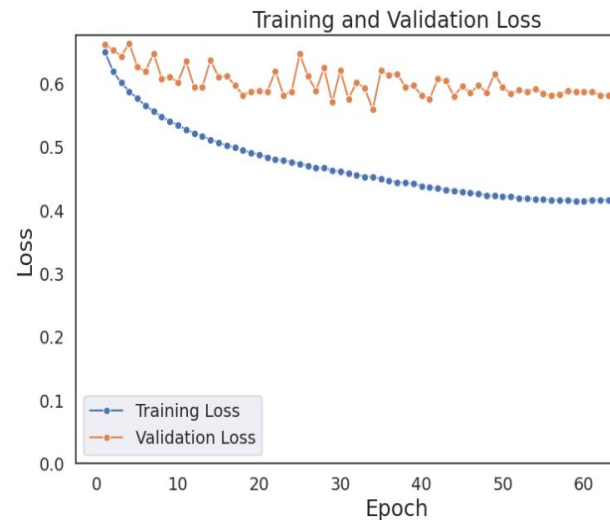
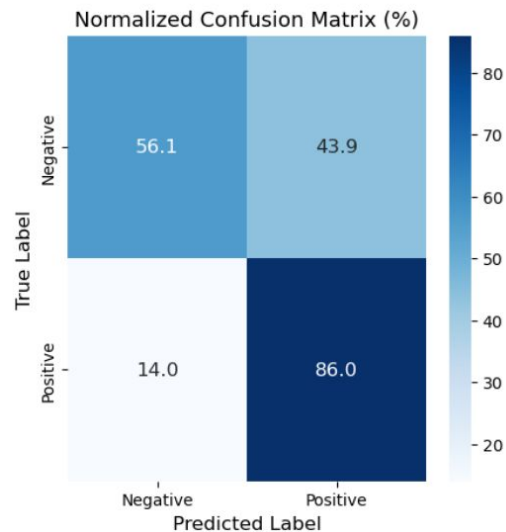
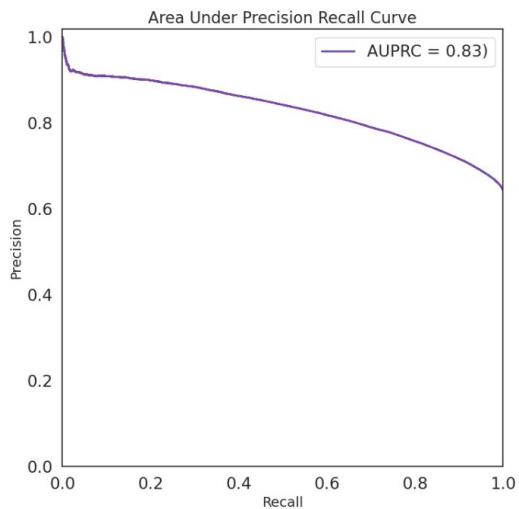


Recall



Precision

Evaluation



Further hypertuning in progress!

Future Developments



Product Focus

Improve model predictions on specific drugs groups (i.e. antibiotics)



Modeling

Graph base GNN
Additional features:
target proteins



Encoding

Train BioBERT with drugs target & action pathway texts



Latency

Inference with GPU instances



Accomplishments

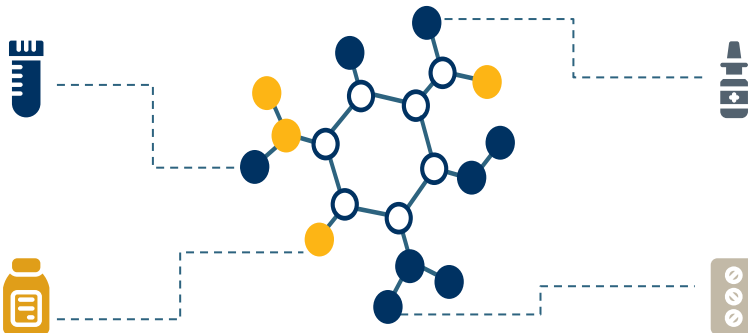


New Drug Development

Inform potential DDI
with drug formula

Scalability

Autoscaling with AKS



NLP Model Serving

Morgan Similarity +
BioClinical BERT

Interpretable Results

Graphs and
downloadable CSV



Mission



**Reimagine drug development by
transforming drug-drug interaction detection**





Q&A
Thank you!

dd-ai-predict.streamlit.app

Acknowledgements

Professors:

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- Mai Do, RPh., MBA
- Dr. Tuan La, PharmD

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2. [Percent of ADE](#)
3. [Adults using multiple medications](#)
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10. [Google/Pegasus-Pubmed](#)
11. [Publicly Available Clinical BERT Embeddings](#)



Thanks

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