



# Kairós

Predicting Drug Interactions

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Matt Kirk  
Elio Qarri  
Michael Jaweed

# Our Team



**Quazi Fairooz**



**Jonathan Tran**



**Matt Kirk**



**Elio Qarri**



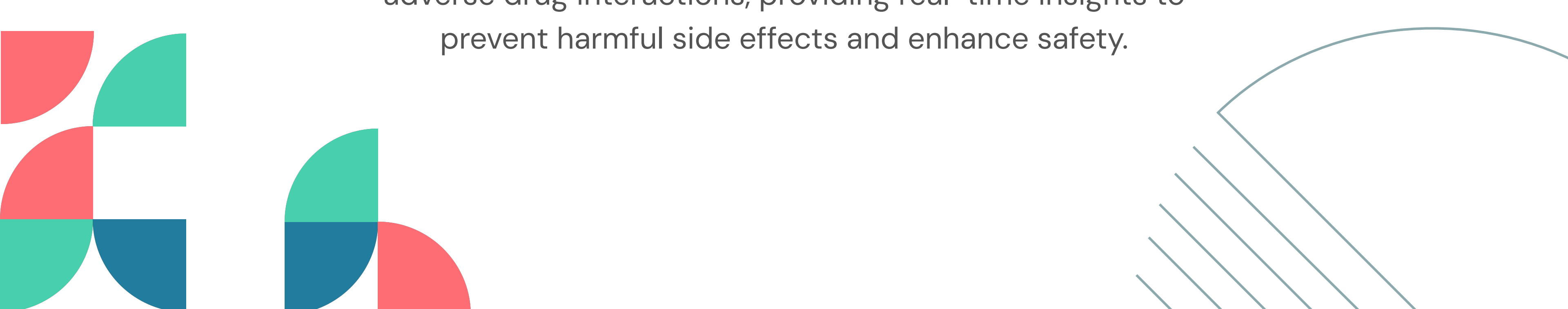
**Michael Jaweed**



# Our Solution



**Kairós** is a web app that uses machine learning to predict adverse drug interactions, providing real-time insights to prevent harmful side effects and enhance safety.

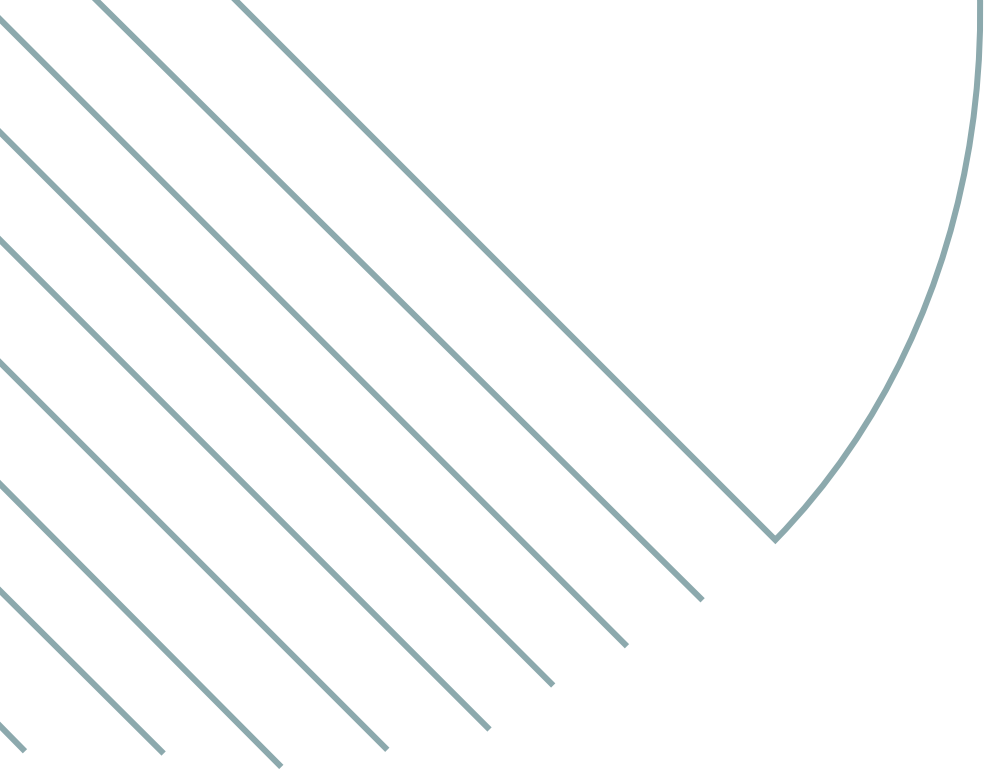




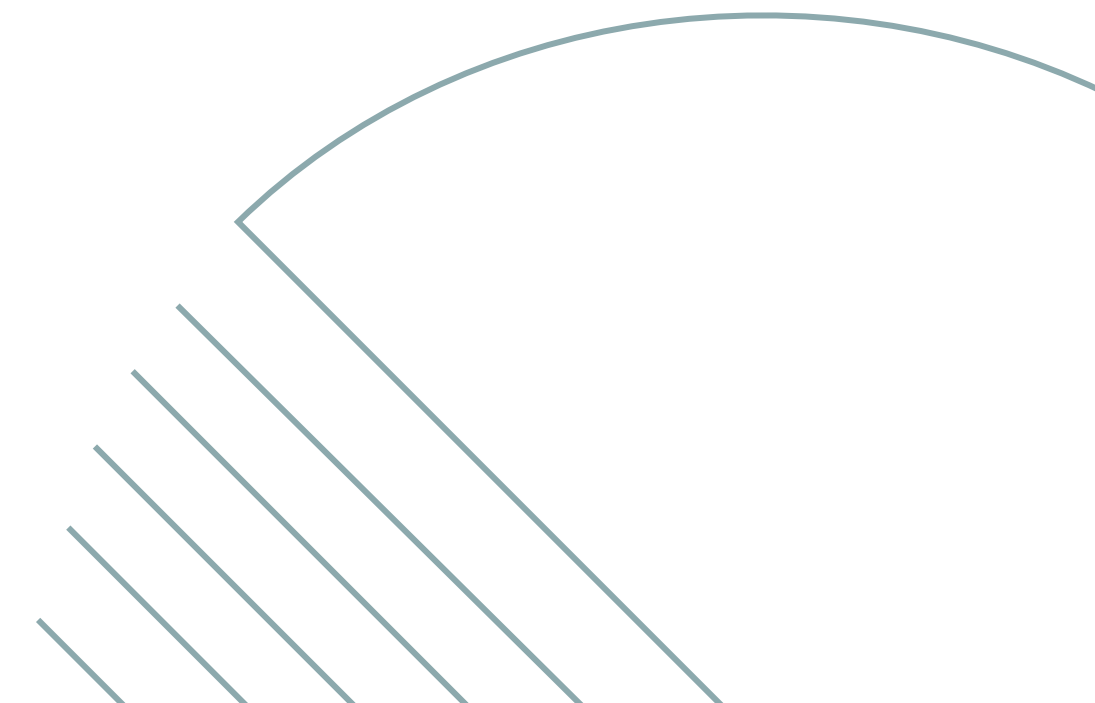
# Why?

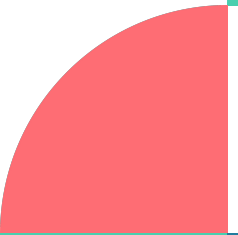
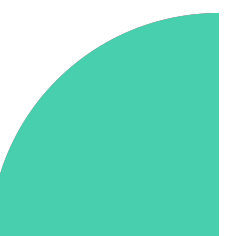
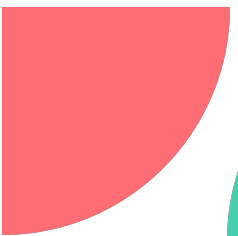
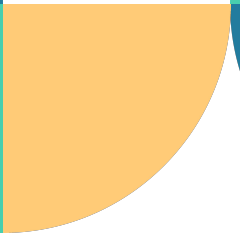
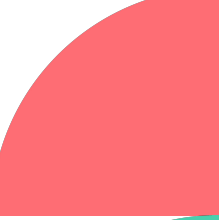
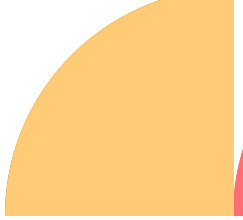
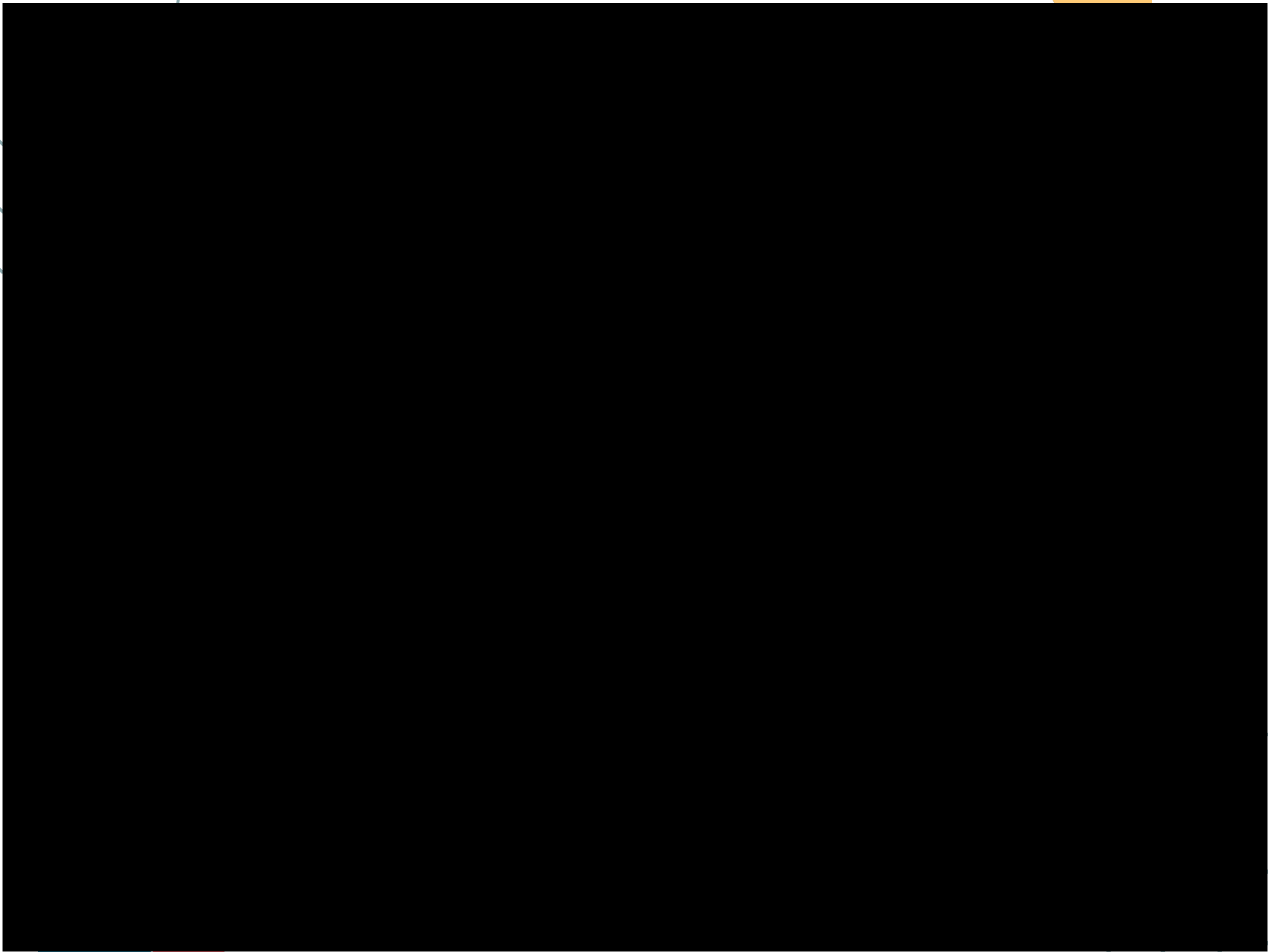
The FDA estimates **2 million serious adverse drug events** and 100,000 deaths annually in the U.S. due to drug interactions.





# MVP Demo

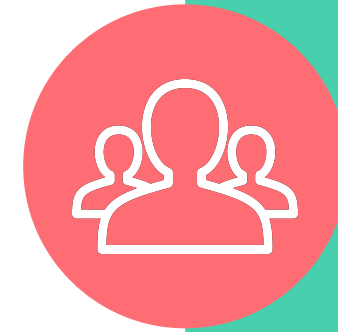




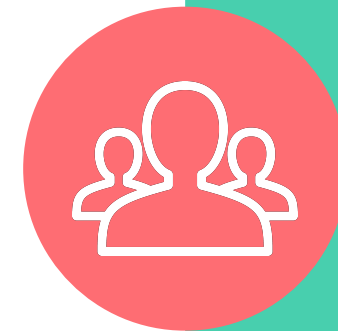


## Target Users

**Individuals taking prescription or over-the-counter medications**



A diabetic person seeks to understand if their current prescription will interact with a new supplement they are considering.



A person wants to determine if a new metabolism booster could adversely affect their health while using a specific antibiotic.

# Market Research

Competitors	Target Users	Uses ML	User Data	OCR
WebMD	Consumers	✘	♥	✘
Drugs.com	Consumers	✘	♥	✘
Medscape	Consumers / Medical Pros	✘	♥	✘
AlphaFold3	Scientists	♥	✘	✘
Kairos	Consumers	♥	♥	♥

Kairós is the only tool that integrates ML modeling, User Data, and OCR into one simple to use platform.





# DATA SOURCES & EDA

**Size:** 192,347 drug-to-drug interaction pairs

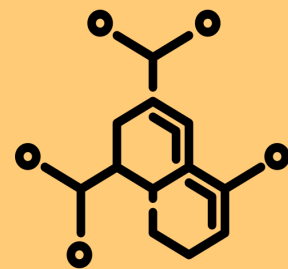
**Split:** Train (70%), Validation (15%), Test (15%)

**Target Variable:** Boolean for:

- Acetylation
- Amidation
- Hydrolysis of Amide



DrugBank



ChEMBL



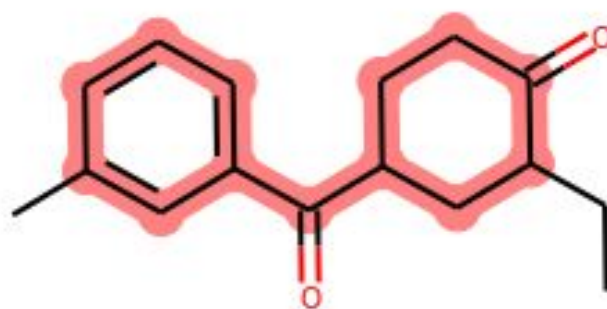
NIH Dietary Supplements

# Reaction Modeling

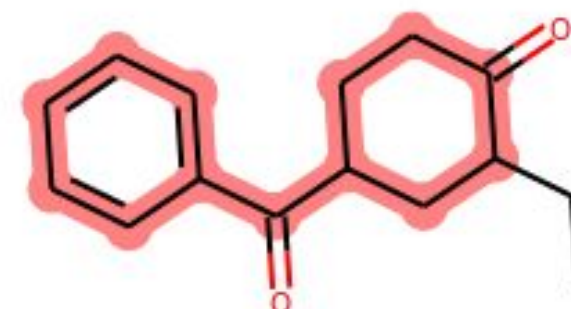
		Canonical Smiles			Canonical Smiles	RDKit Reaction	
Drug1_ID		Drug1	drug1_name	Drug2_ID	Drug2	drug2_name	acetylation_reaction_flag
0	DB00855	NCC(=O)CCC(O)=O	Aminolevulinic acid	DB00460	COC(=O)CCC1=C2NC(\C=C3/N=C(/C=C4\N\C(=C/C5=N/C...	Verteporfin	1
1	DB09536	O=[Ti]=O	Titanium dioxide	DB00460	COC(=O)CCC1=C2NC(\C=C3/N=C(/C=C4\N\C(=C/C5=N/C...	Verteporfin	0
2	DB09536	O=[Ti]=O	Titanium dioxide	DB00460	COC(=O)CCC1=C2NC(\C=C3/N=C(/C=C4\N\C(=C/C5=N/C...	Verteporfin	0
3	DB01600	CC(C(O)=O)C1=CC=C(S1)C(=O)C1=CC=CC=C1	Tiaprofenic acid	DB00460	COC(=O)CCC1=C2NC(\C=C3/N=C(/C=C4\N\C(=C/C5=N/C...	Verteporfin	1
4	DB09000	CC(CN(C)C)CN1C2=CC=CC=C2SC2=C1C=C(C=C2)C#N	Cyamemazine	DB00460	COC(=O)CCC1=C2NC(\C=C3/N=C(/C=C4\N\C(=C/C5=N/C...	Verteporfin	0
...	...	...	...	...	...	...	...
185402	DB00281	CCN(CC)CC(=O)NC1=C(C)C=CC=C1C	Lidocaine	DB06708	CCCCN(CCCC)CC(O)C1=C2C(=CC(Cl)=C1)\C(=C/C1=CC=...	Lumefantrine	1
185403	DB01088	[H][C@]12C[C@@H](O)[C@H](\C=C\C[C@@H](O)C(C)CC#...	Iloprost	DB01235	N[C@@H](CC1=CC(O)=C(O)C=C1)C(O)=O	Levodopa	1
185404	DB00857	CN(C\C=C\C#CC(C)(C)C)CC1=CC=CC2=CC=CC=C12	Terbinafine	DB00196	OC(CN1C=NC=N1)(CN1C=NC=N1)C1=C(F)C=C(F)C=C1	Fluconazole	0
185405	DB00734	CC1=C(CCN2CCC(CC2)C2=NOC3=C2C=CC(F)=C3)C(=O)N2...	Risperidone	DB02703	[H][C@@]12C[C@@H](O)[C@@]3([H])[C@@]4(C)CC[C@@...	Fusidic acid	0
185406	DB00356	ClC1=CC2=C(OC(=O)N2)C=C1	Chlorzoxazone	DB00934	CNCCCC12CCC(C3=CC=CC=C13)C1=CC=CC=C21	Maprotiline	0



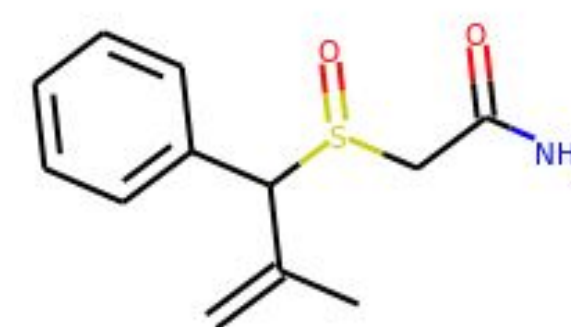
Open-Source Cheminformatics  
and Machine Learning



c1ccc(CC2CCCCC2)cc1



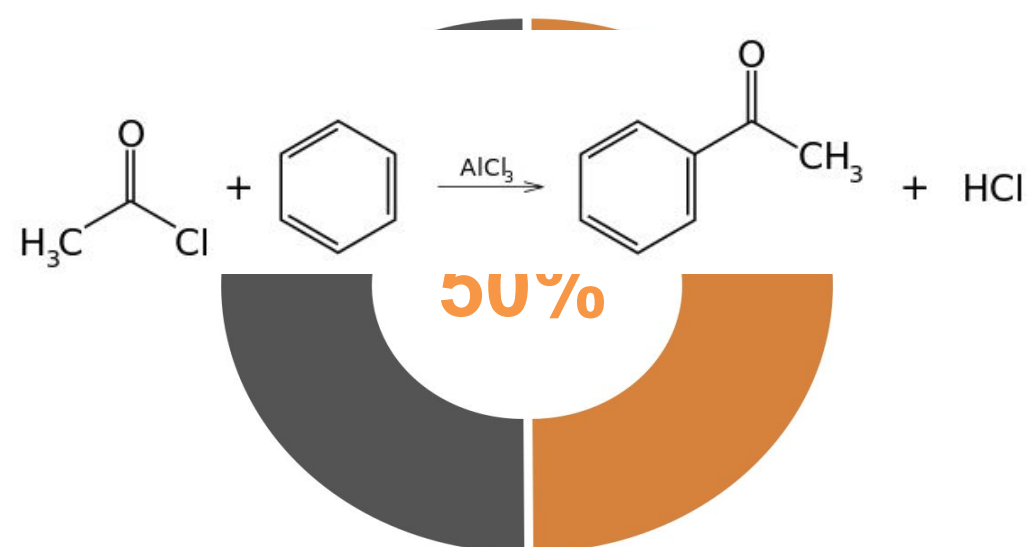
c1ccc(CC2CCCCC2)cc1



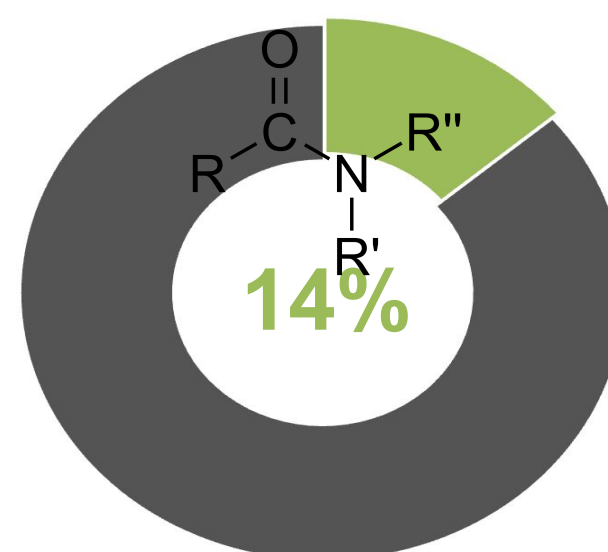
c1ccccc1

# Type of Reactions

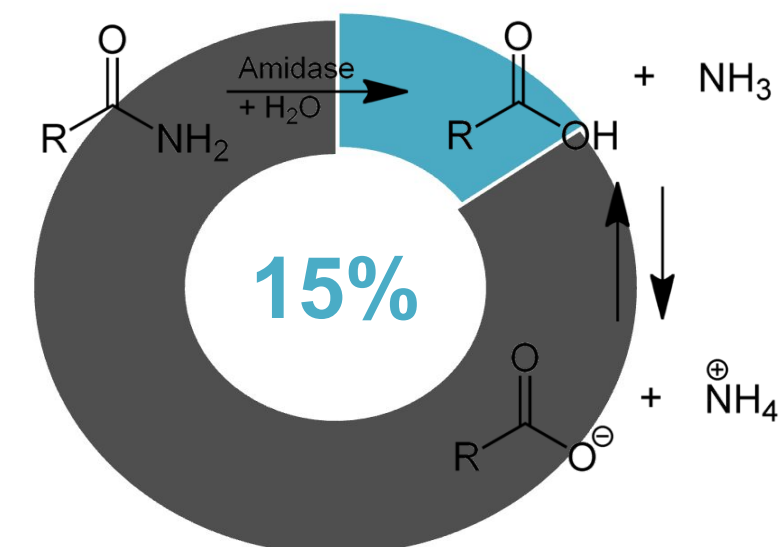
## Acetylation



## Amidation



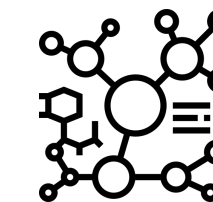
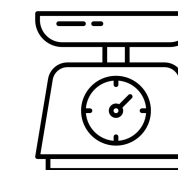
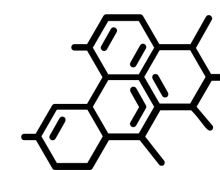
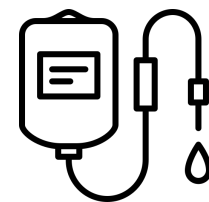
## Hydrolysis of Amide



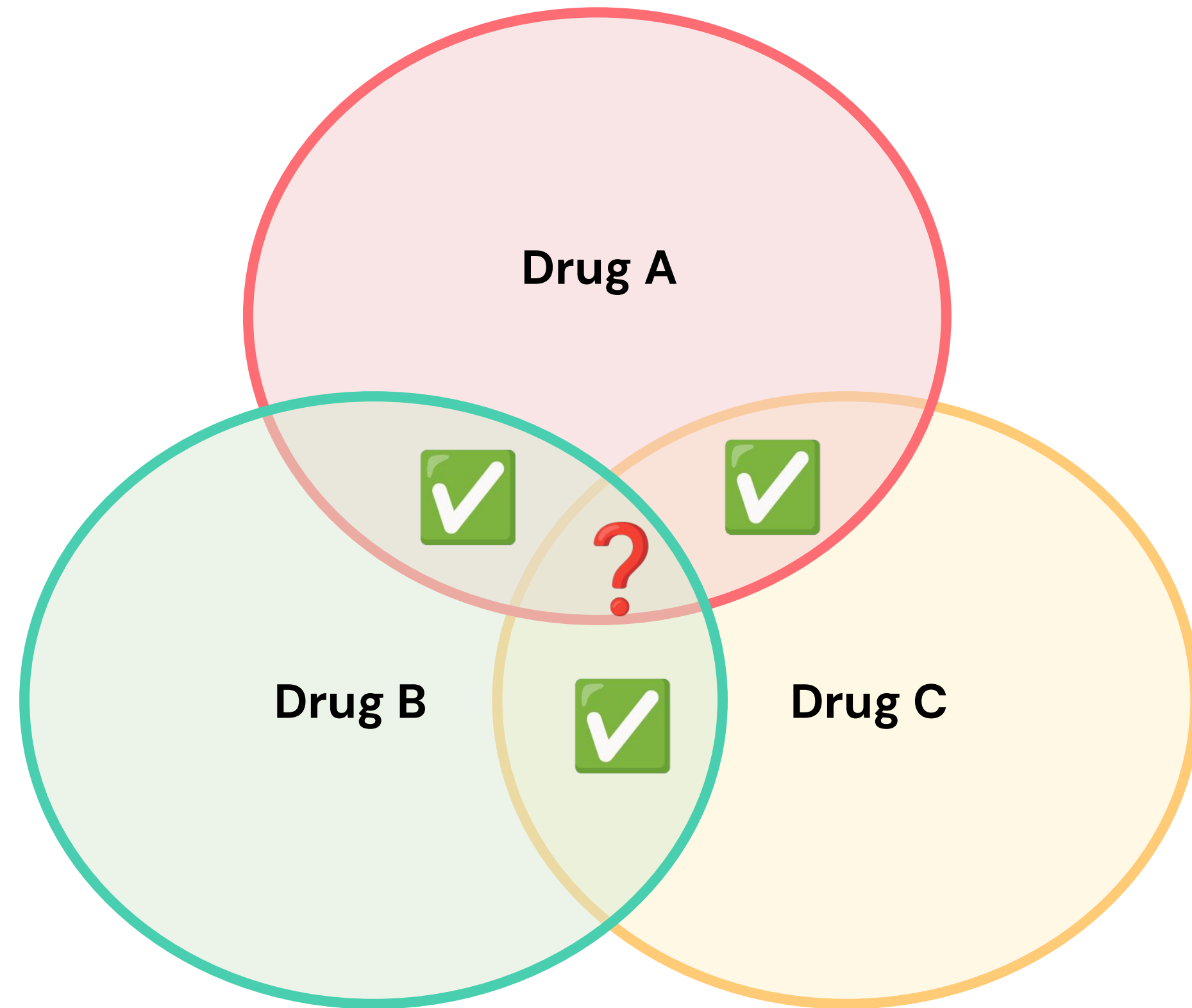
Each reaction is generated with a different reaction template, resulting in 3 sets of target variables with different class distributions

# Feature Selection

		Route of Administration			Molecular Properties			Hydrogen Bond Acceptor / Donor	
	name	oral	parenteral	topical	alogp	aromatic_rings	full_molecular_weight	hba	hbd
0	LISINOPRIL ANHYDROUS	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	1.24	1	405.5	5	4
1	NAPROXEN	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	3.04	2	230.26	2	1



# Model Limitation: Multiple Drug Interactions



Guided screen for synergistic three-drug combinations:<https://www.ncbi.nlm.nih.gov/pmc/articles/PMC7347197/>

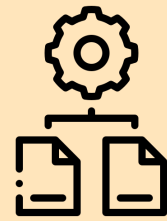
Design of high-order antibiotic combinations against *M. tuberculosis* by ranking and exclusion:<https://www.nature.com/articles/s41598-019-48410-y>

Systematic exploration of synergistic drug pairs:<https://www.ncbi.nlm.nih.gov/pmc/articles/PMC3261710/>

# ML PIPELINE

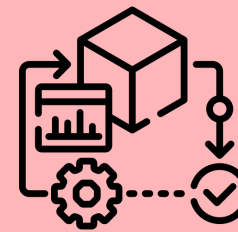
## DATA PREP

- Dataset Ingestion
- Data Preprocessing
- Feature Engineering / Selection



## ML MODELING

- Model Selection
- Model Training
- Hyperparameter Tuning



## MODEL TESTING

- Model Evaluation
- Model Selection
- Addl. Training & Data Acquisition

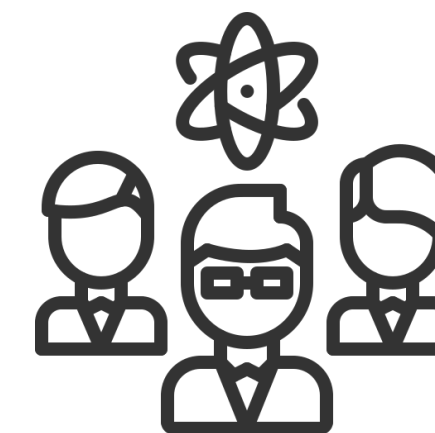
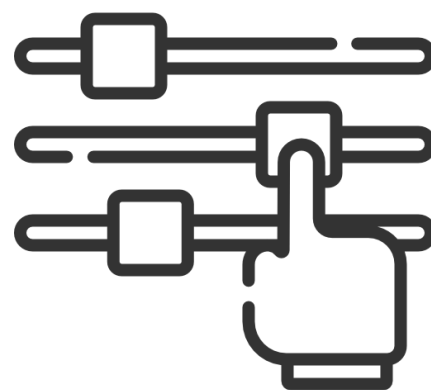
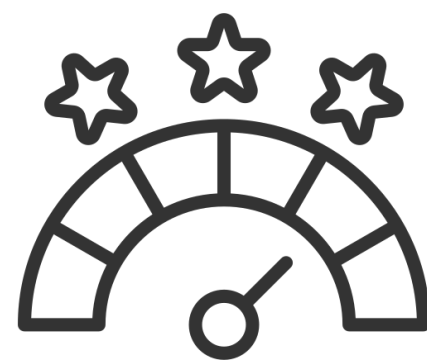


# Model Results

Reaction	Model	Mean F2 Score		Mean F1 Score		Mean PR AUC	
		Val	Train	Val	Train	Val	Train
Acetylation	XGBoost	0.995	0.996	0.996	0.997	0.997	0.999
	Random Forest	0.973	0.999	0.965	0.999	0.971	0.999
	SVM	0.799	0.803	0.789	0.793	0.839	0.842
	Logistic Regression	0.620	0.611	0.643	0.633	0.747	0.738
Amidation	XGBoost	0.988	0.990	0.983	0.990	0.989	0.998
	Random Forest	0.954	0.949	0.939	0.935	0.982	0.980
Hydrolysis of Amide	XGBoost	0.993	0.989	0.993	0.990	0.993	0.997
	Random Forest	0.853	0.827	0.88	0.862	0.903	0.951

# Best Models vs Test Data

Reaction	Model	Mean F2 Score			Mean F1 Score			Mean PR AUC		
		Test	Val	Train	Test	Val	Train	Test	Val	Train
Acetylation	XGBoost	0.996	0.995	0.996	0.996	0.996	0.997	0.997	0.997	0.999
Amidation	XGBoost	0.990	0.988	0.990	0.991	0.983	0.990	0.992	0.989	0.998
Hydrolysis of Amide	XGBoost	0.992	0.993	0.989	0.992	0.993	0.990	0.993	0.993	0.997





# Future Enhancements & Goals

## User Data Management & Security

- Expand User Profile Information
- Personalized User Experience
- Use strict user data privacy & protocols

## Chatbot & OCR Improvements

- Improve Chatbot experience
- Include other languages for chatbot and OCR

## Reaction Severity Score

- Integrate more datasets in order to predict severity score
- Add High / Medium / Low Score
- Update the ML models

# Kairós

Multiple Drug Interactions

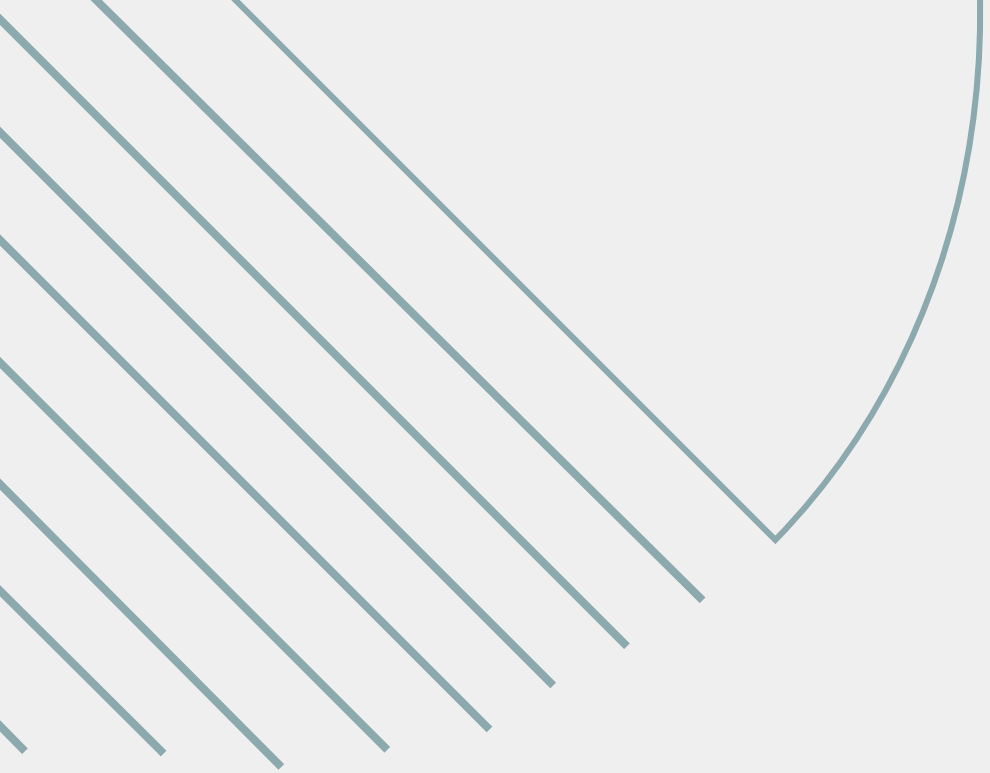
ML Modeling Integrated

User Input / Login

OCR Integrated

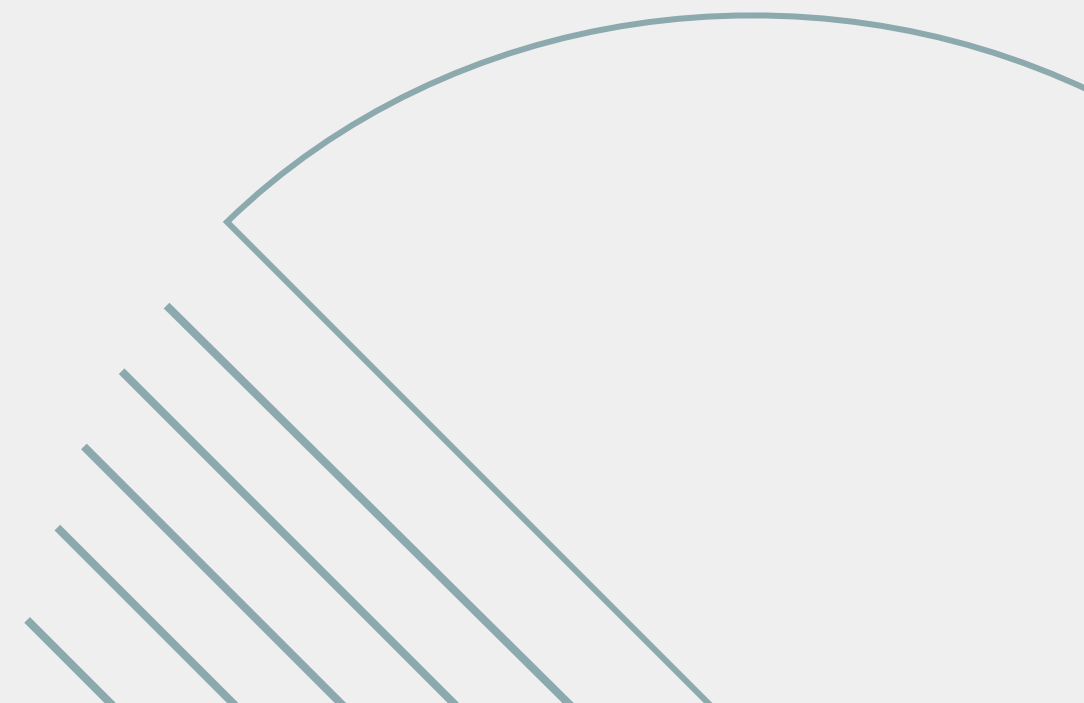
3 Types of Reactions





# THANK YOU

Kairós team

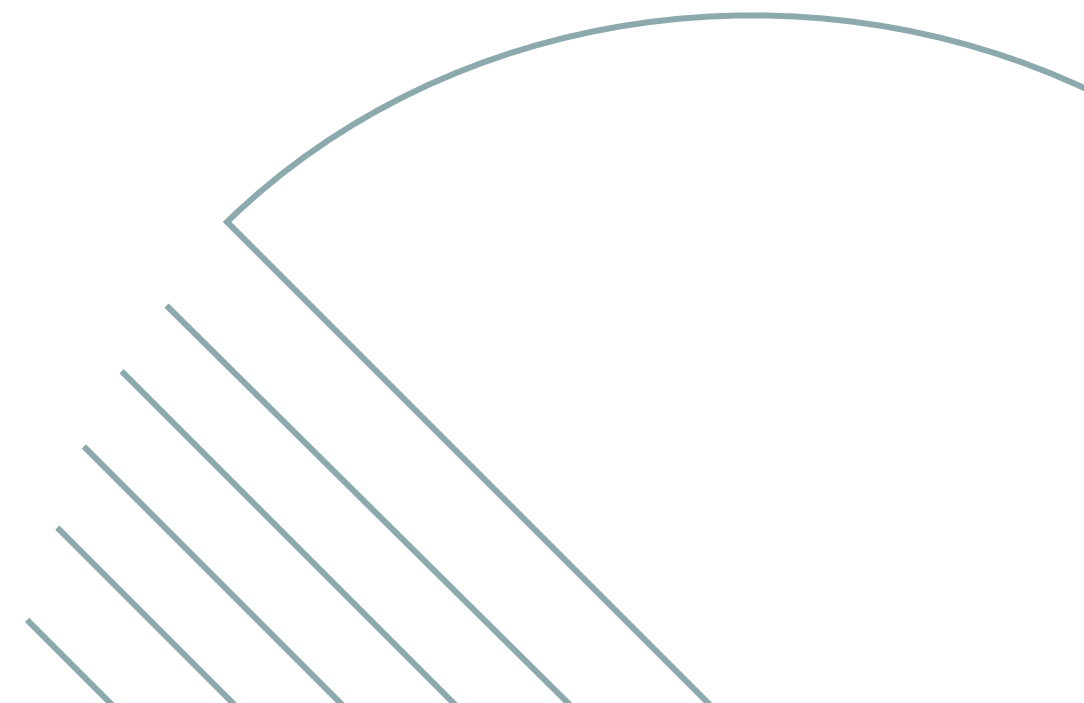
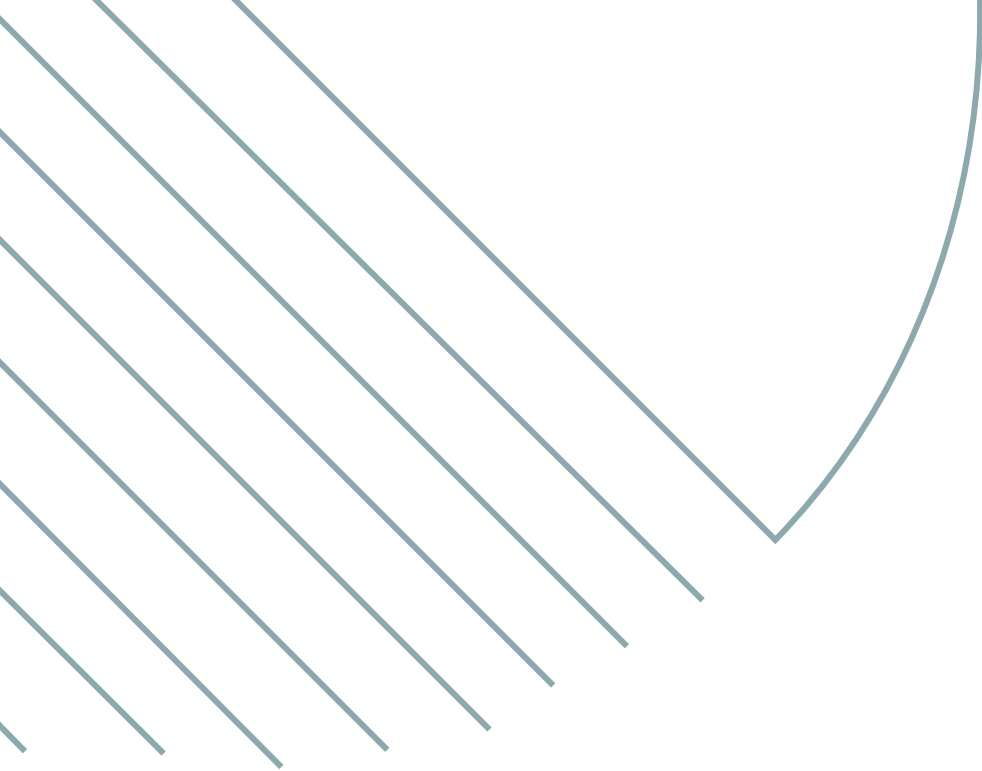


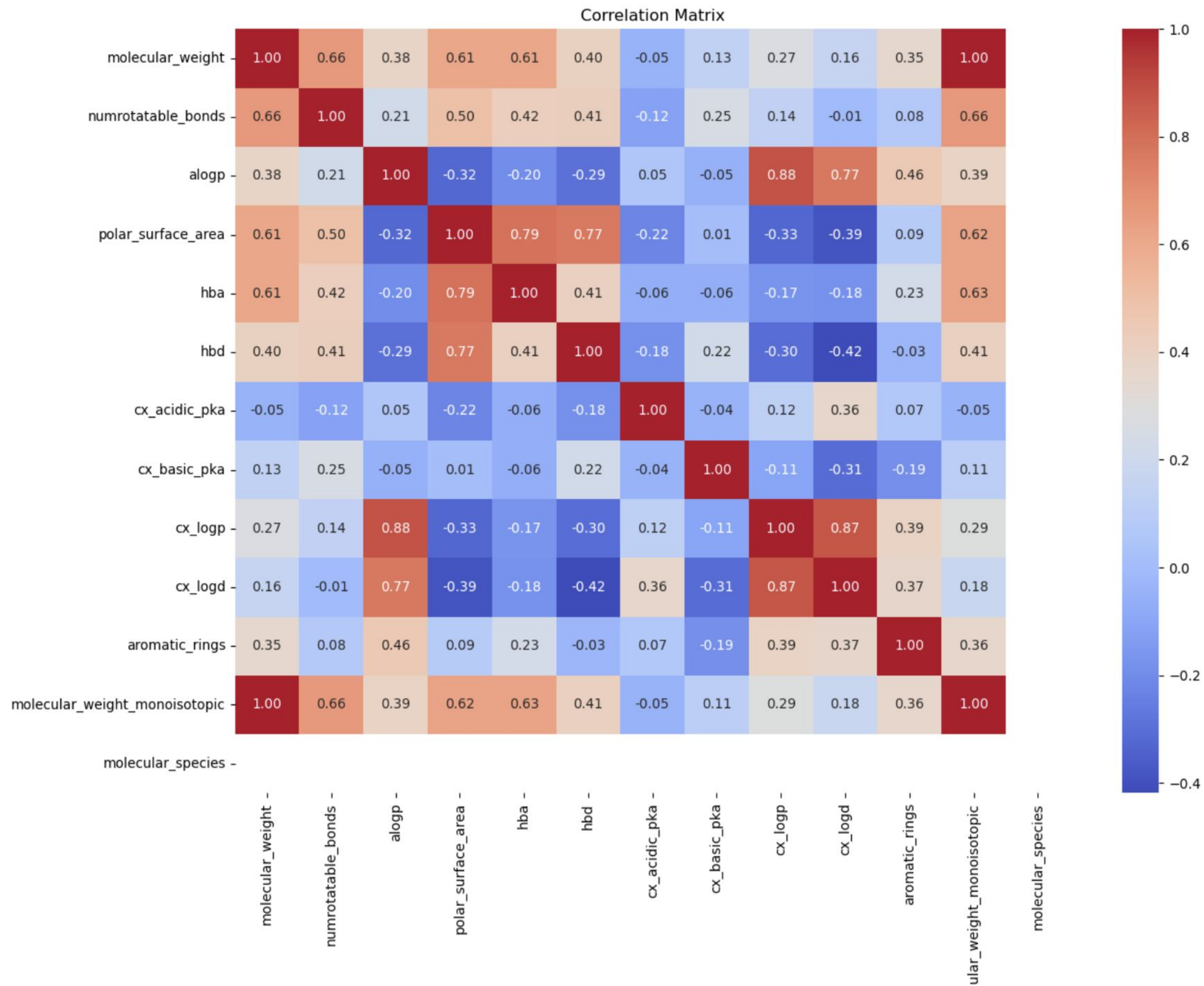


# SUMMARY

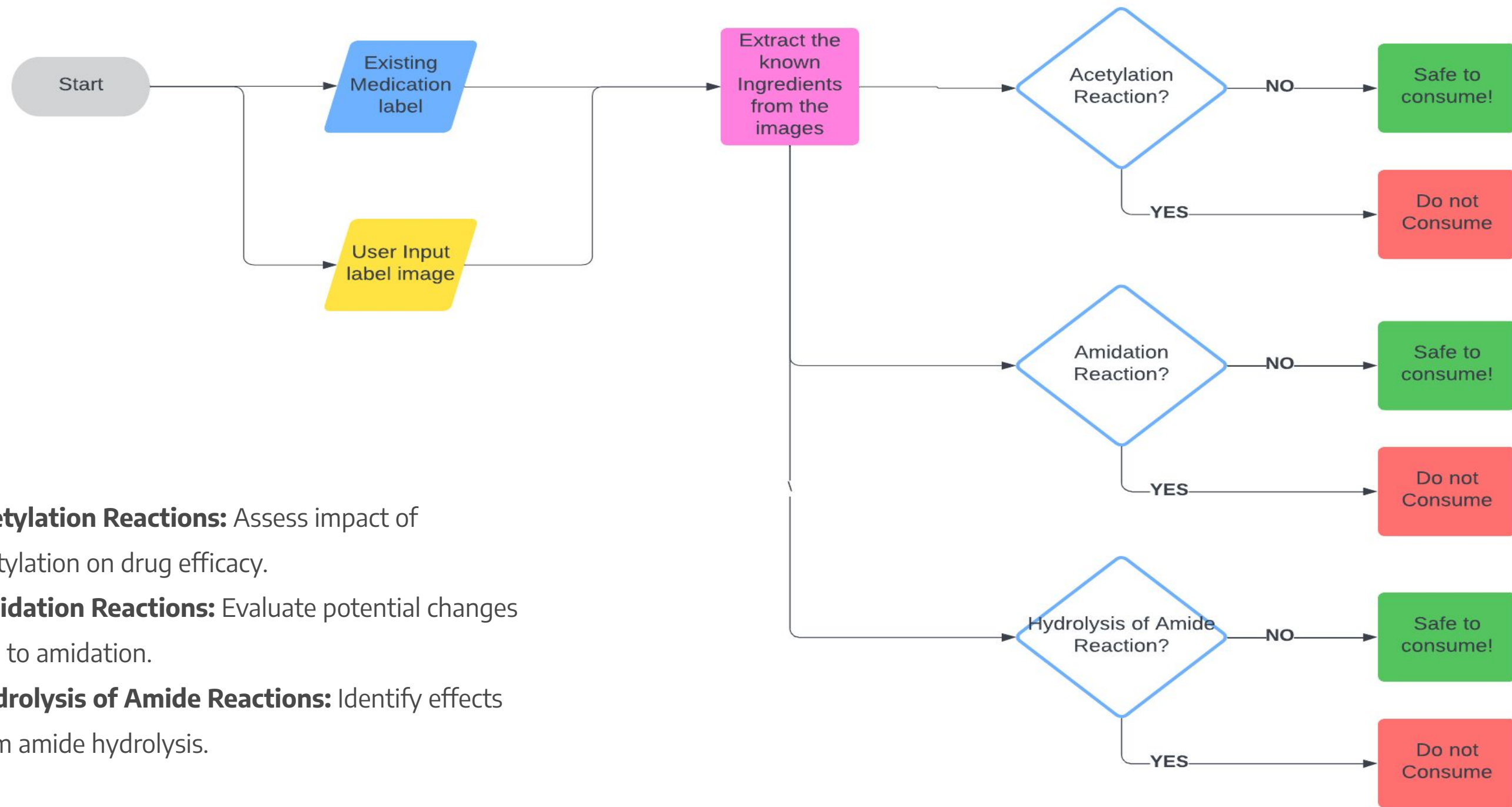
Lorem ipsum dolor sit amet, consectetur adipiscing elit. Etiam mattis, nunc vitae eleifend posuere, turpis mauris vestibulum purus, in pellentesque tellus elit vel nisl. Nam elementum nunc quis sapien pretium, at tincidunt mauris dignissim.

# Appendix





# MVP Flow



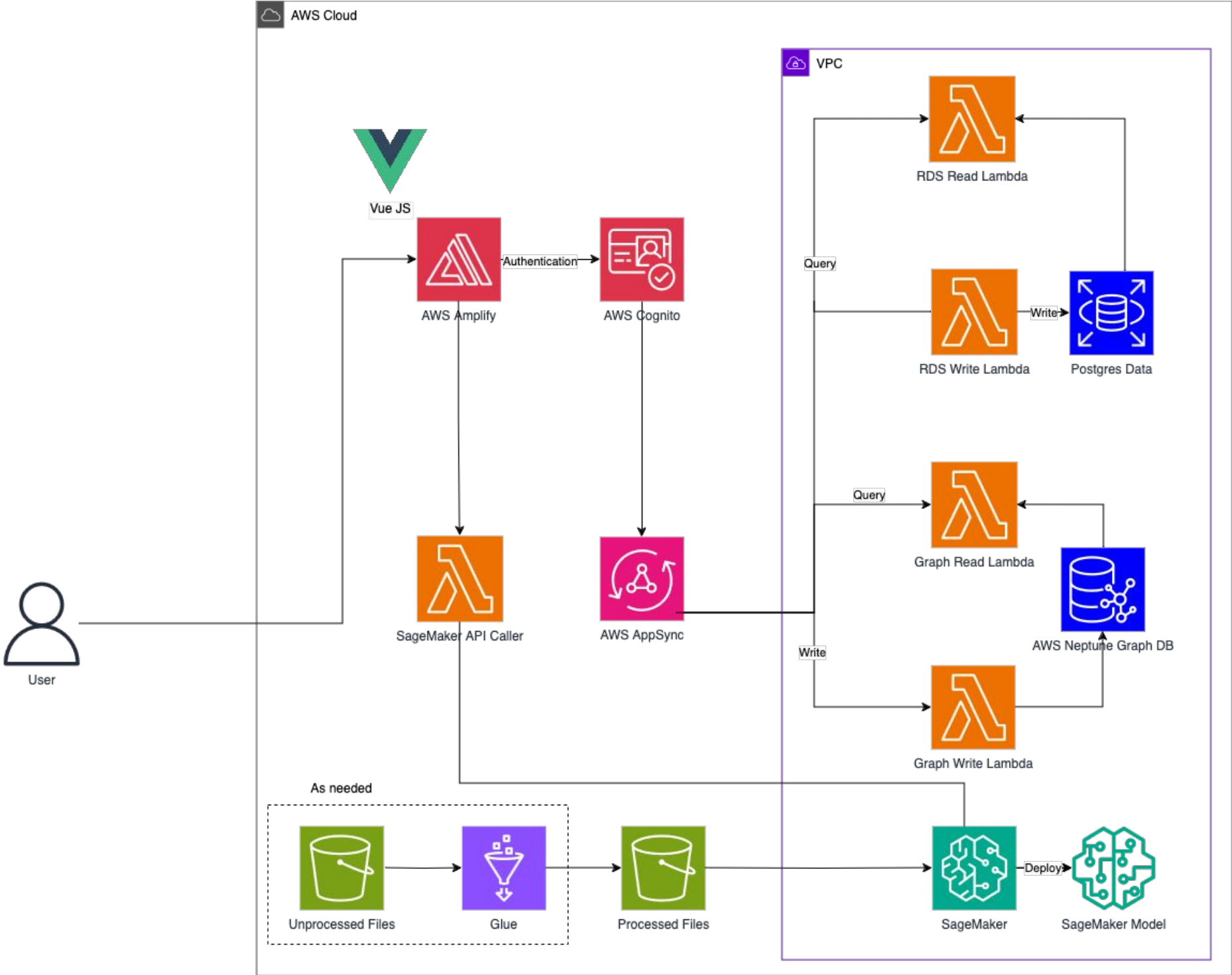
**Acetylation Reactions:** Assess impact of acetylation on drug efficacy.

**Amidation Reactions:** Evaluate potential changes due to amidation.

**Hydrolysis of Amide Reactions:** Identify effects from amide hydrolysis.



# Data Architecture






# OCR Integration

Limit 200MB per file • JPG, JPEG, PNG

IMG\_4902.JPG 1.3MB



Uploaded Image Thumbnail

Scan Medications

Potential Medications from Image:

Select a medication to add:

TAMSULOSIN

Add Selected Medication

Enter a drug name

VANOXERINE

Search Drug

Select the correct drug:

VANOXERINE

Add Selected Drug

TAMSULOSIN



Amazon Textract

# Type of Reactions

## Acetylation

Many drugs are processed in the body through the acetylation reaction, either by biotransformation into an effective compound or to be metabolized into substances that the body can excrete in a more simpler manner.

## Amidation

Amides are polar, meaning they have regions of high positive and negative electrical charge density, which allows them to interact with biological receptors and enzymes. Amides are also stable and can help drugs resist rapid metabolic degradation in the human body.

## Hydrolysis of Amide

Amide hydrolysis is important in drug discovery and development because it can make drugs active

