



Fingerprint-Based Machine Learning Prediction of Chemical Properties

Presented by

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First Semester

Introduction

Background, Object and Scope

Selection of suitable substances for production processes is a long-standing problem in the petrochemical industry. Development of a model by using machine learning (ML) become more popular because they are faster and more accurate than the traditional methods



Fig 1. Chemical Experiment

Objective

1. To study the converting of SMILE structures into molecular fingerprints.
2. To model and improve the accurate of prediction model on T_b and P^{sat} of substances from molecular fingerprints using machine learning and techniques.

Scope

1. To study T_b and P^{sat} of pure organic compound that contain C, H, O and N atom and Number of C atom is 1-12 atom
2. To study morgan fingerprint that one of tool in RDKit Library using Python Programming Language

Introduction

Review Previous Work

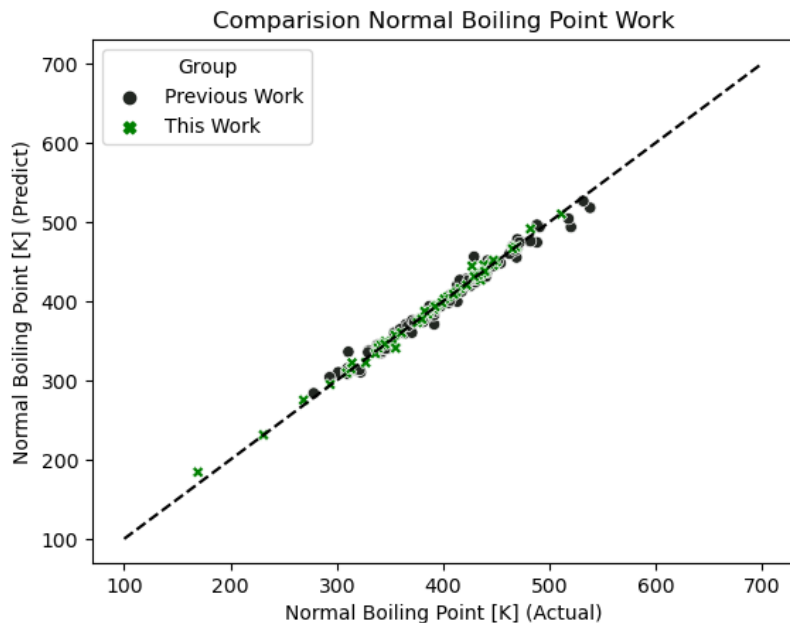
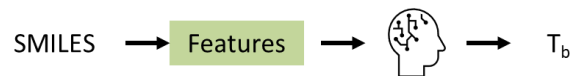


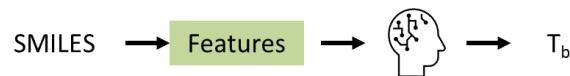
Table 1. Performance of Each Work on T_b Prediction Model within CH Scope

Error Metric	Previous Work [2]	This Work
MAE	5.86	3.20
MAPE(%)	1.47	0.84
RMSE	7.92	4.70
R^2	0.98	0.99

Fig 2. Comparison T_b Work on CH Scope

Introduction

Review Previous Work



The **Previous work** (Nattasinee and colleagues) ^[2] use Machine Learning (ML) by convert the representation of molecule from text format (SMILES) into numbers in the following table while **This work** convert SMILES to “Molecular Fingerprint”

Table 2. Compare Prediction Between Each Work of Similar Structure Substance

SMILES	C	Double	Triple	Branch	Cyclic	Actual T _b	Predict T _b Previous Work ^[2]	Predict T _b This Work
C1CCC=CCC1	7	1	0	0	1	388.15	375.85	387.34
CC1=CCCCC1	7	1	0	0	1	383.45	375.85	377.95
CC1CCC=CC1	7	1	0	0	1	375.85	375.85	370.31

Similar
SMILES

Same
Features

Same
Predictions

Introduction

Result

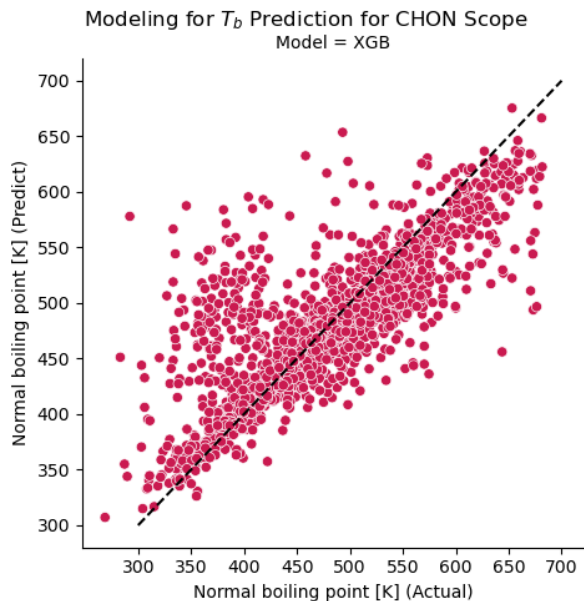
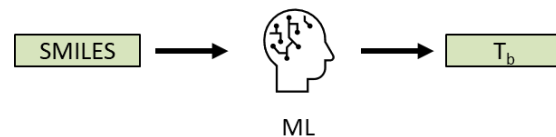


Table 3. Performance on T_b Prediction Model within CHON Scope

Dataset	MAE	MAPE (%)	RMSE	R^2
Train	20.42	4.51	30.38	0.83
Test	26.72	5.94	43.40	0.66

Fig 3. Best T_b model on CHON Scope

Introduction Result

T_b Model Prediction for CHON Scope

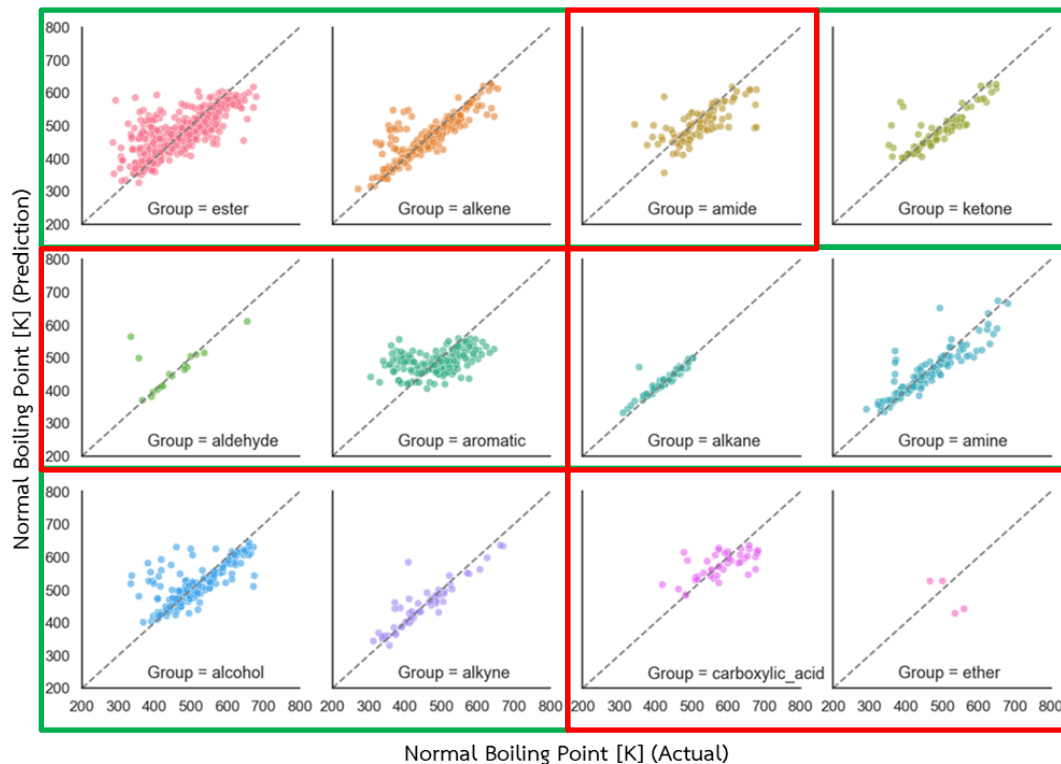
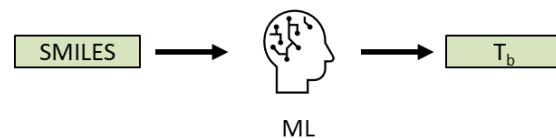


Fig 4. T_b Prediction on CHON Scope Group by Functional Group



The functional group that this T_b prediction model can predict well is that the group that have condition with $R^2 \geq 0.5$ and $MAPE \leq 10\%$

Acceptable: Ester, Alkene, Ketone, Alkane, Amine, Alcohol, Alkyne

Unacceptable: Amide, Aldehyde, Aromatic, Carboxylic Acid, Ether

Second Semester

Methodology

Data Collection

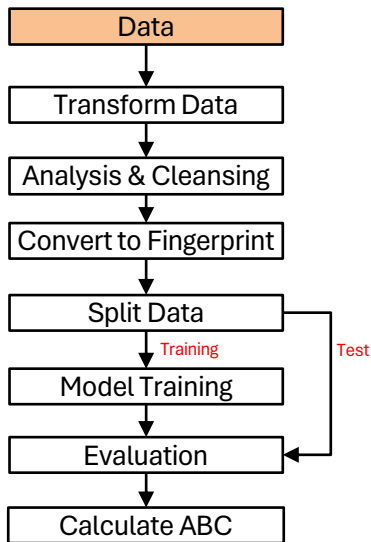
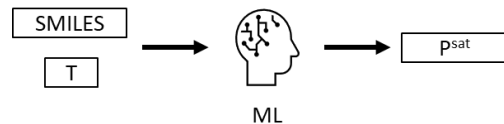


Table 4. ChEDL Database Collection

Database Source	Datapoint
Hall, K. R. Vapor Pressure and Antoine Constants for Hydrocarbons, and S, Se, Te, and Halogen Containing Organic Compounds. Springer, 1999.	6,346
Dykyj, J., and K. R. Hall. "Vapor Pressure and Antoine Constants for Oxygen Containing Organic Compounds". 2000.	
Hall, K. R. Vapor Pressure and Antoine Constants for Nitrogen Containing Organic Compounds. Springer, 2001.	

The Total amount of Substance that be included in CHON is 2,246 datapoints.

Methodology

Transform Data

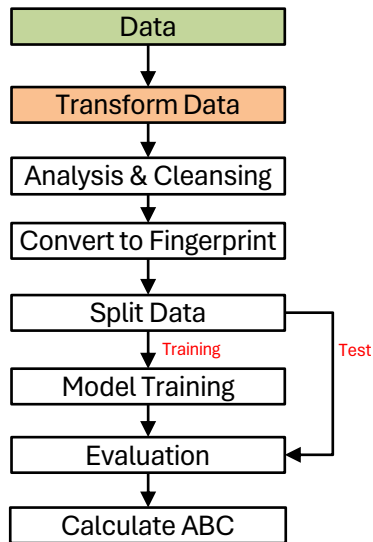
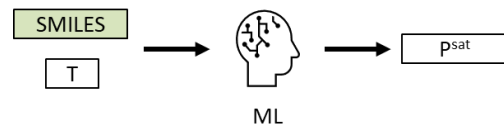


Table 5. P^{sat} Raw data

CAS	Name	A	B	C	Tmin	Tmax
50-00-0	Methanal	21.37	2204.13	-30.15	190	271
51-66-1	p-Methoxy-acetanilide	20.27	3916.19	-177.10	456	533
51-75-2	N-Methylbis(2-chloroethyl)amine	25.61	6563.29	0	273	333
541-35-5	Butyramide	22.08	4164.35	-109.32	320	398

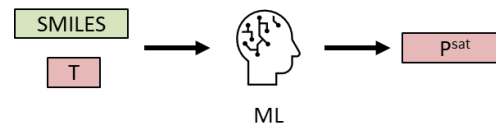


Use PubChem Database to
convert Name of compound to its SMILES

Name	SMILES	A	B	C	Tmin	Tmax
Butyramide	<chem>CCCC(=O)N</chem>	22.08	4164.35	-109.32	320	398

Methodology

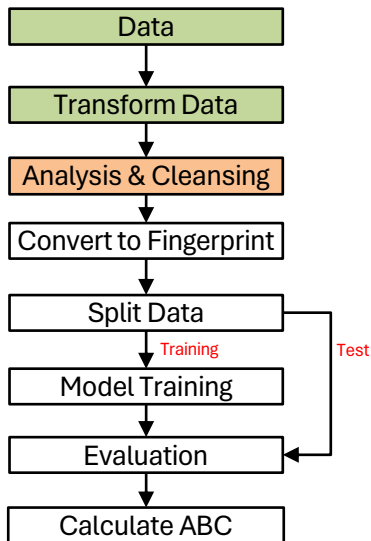
Analysis & Cleansing



Calculate the Vapor Pressure value using the Antoine equation, which is divided into 5 points according to T_{min} and T_{max} range, which can be shown in the table below.

SMILES	A	B	C	T_{min}	T_{max}
CCCC(=O)N	22.08	4164.35	-109.32	320	398

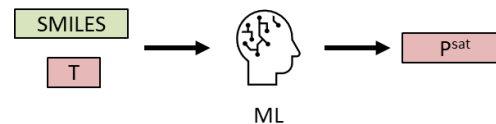
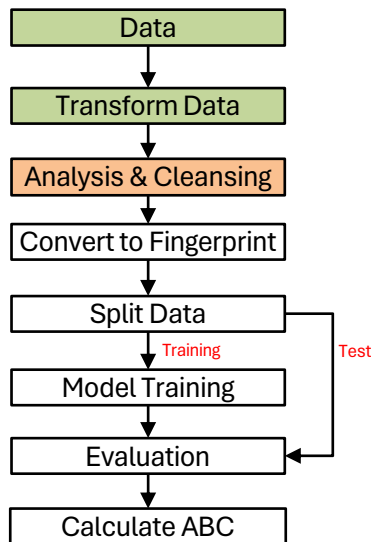
SMILES	VP1	VP2	VP3	VP4	VP5
CCCC(=O)N	2.32	3.99	5.40	6.61	7.66



Note : VP – Vapor Pressure in $\ln(P^{sat})$, P^{sat} in Pa, T in K

Methodology

Analysis & Cleansing



After get the Vapor Pressure value of all substance, use Boxplot to see distribution of Vapor Pressure so that can we detect abnormal Vapor Pressure

Table 6. SMILES and Vapor Pressure Data

SMILES	VP1	VP2	VP3	VP4	VP5
CCCC(=O)N	2.32	3.99	5.40	6.61	7.66
CC(CO)O	9.89	10.36	10.79	11.19	11.57
.
COC1=CC=C(C=C1)CC=C	4.85	7.15	8.93	10.35	11.52

2,246
datapoint

Use boxplot to detect outliers

Outliers

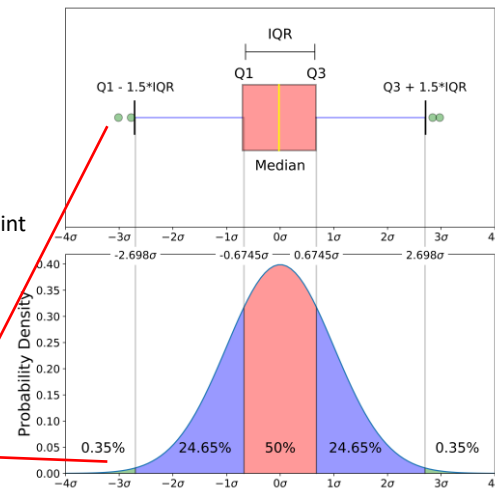
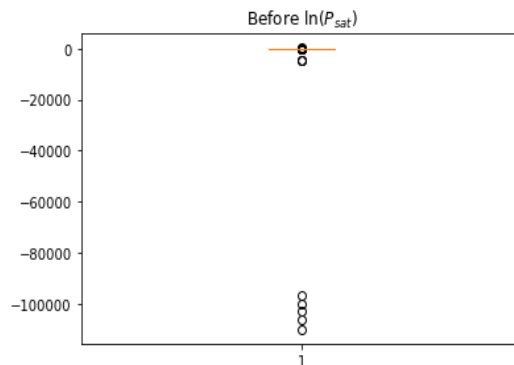
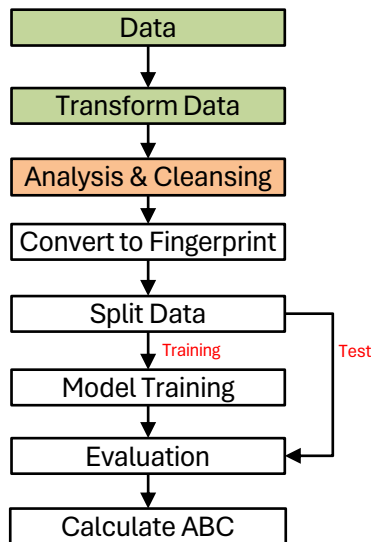


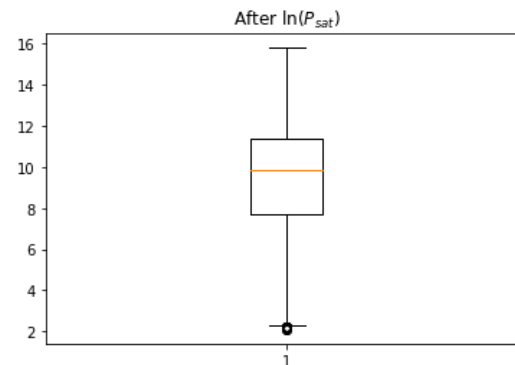
Figure 5. Boxplot and Normal Distribution

Methodology

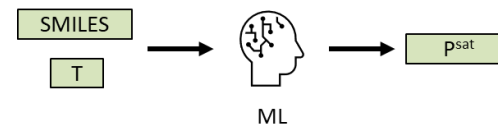
Analysis & Cleansing



Value	$\ln(P^{sat})$
Min	-1.10×10^5
Max	340.72
Point	2,246

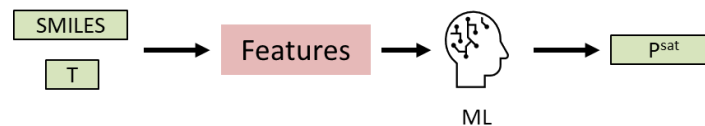
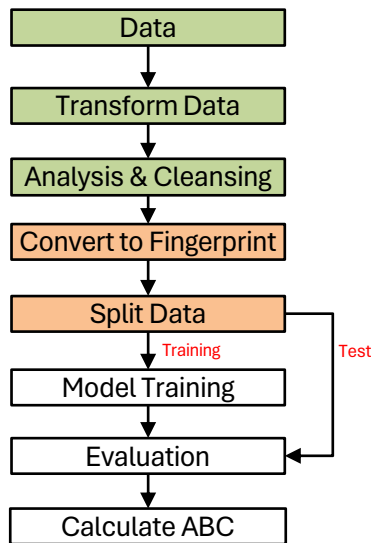


Value	$\ln(P^{sat})$	$P^{sat} \text{ (atm)}$
Min	2.09	8.1×10^{-5}
Max	15.78	71.58
Point	1,787	



Methodology

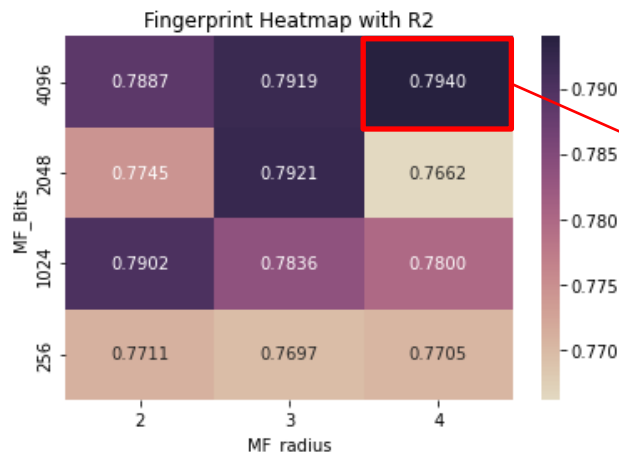
Fingerprint Selecting



All Data		
Alcohol	Alkyne	Carboxylic Acid
Aldehyde	Amide	Ester
Alkane	Amine	Ether
Alkene	Aromatic	Ketone



Training Data	Test Data
80%	20%



The best radius and bits for Fingerprint

Fig 6. Heatmap Fingerprint for P^{sat} modeling

Methodology Training

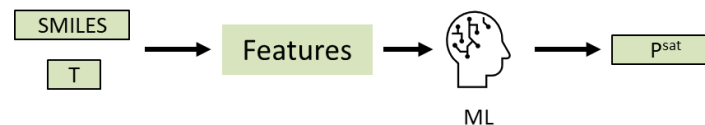
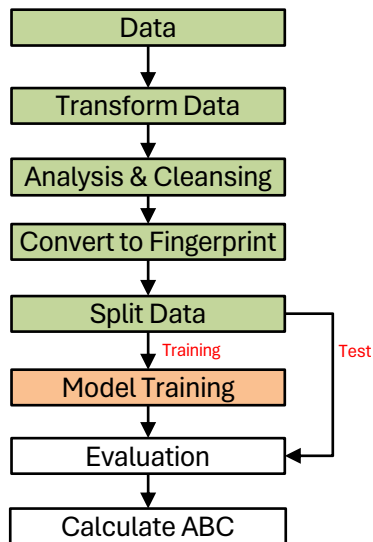


Table 7. Setting of All Learning Algorithms

ML Algorithm	Hyperparameter
DT	max depth = None, min samples leaf = 1, min samples split = 2
RF	max depth = None, max feature = None, n estimators = 200
XGB	max depth = 5, learning rate = None, n estimators = 400
KNN	Algorithm = ball tree, n neighbors = 5, weights = distance

Note : Each ML algorithm use K-Fold with K = 5 and Grid Search

Result Evaluation

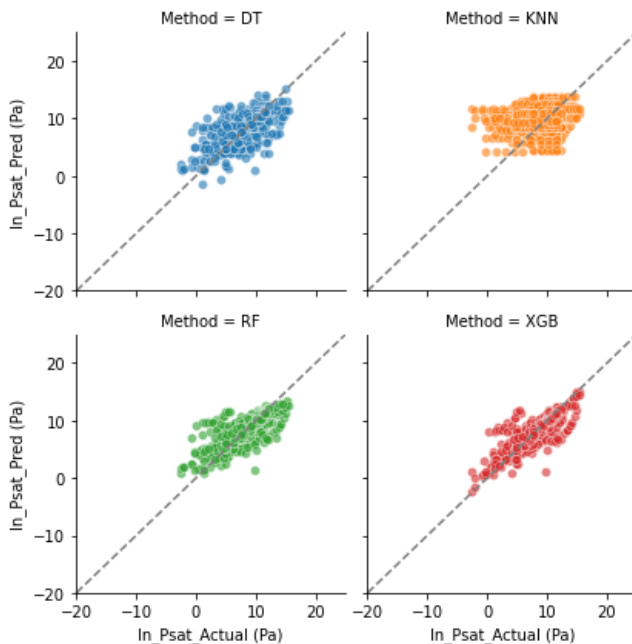
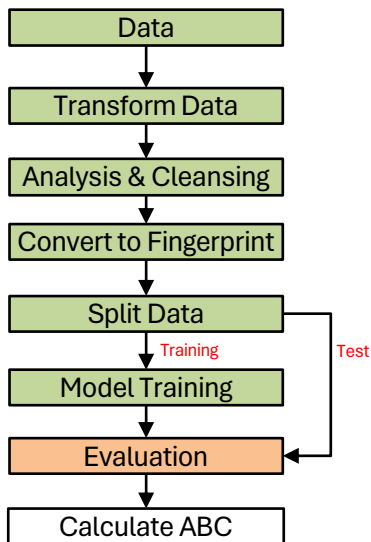


Fig 7. Result of each algorithm

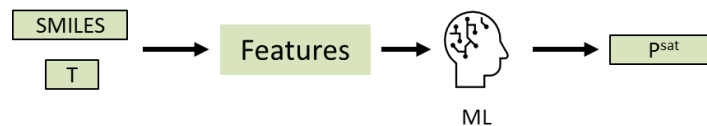
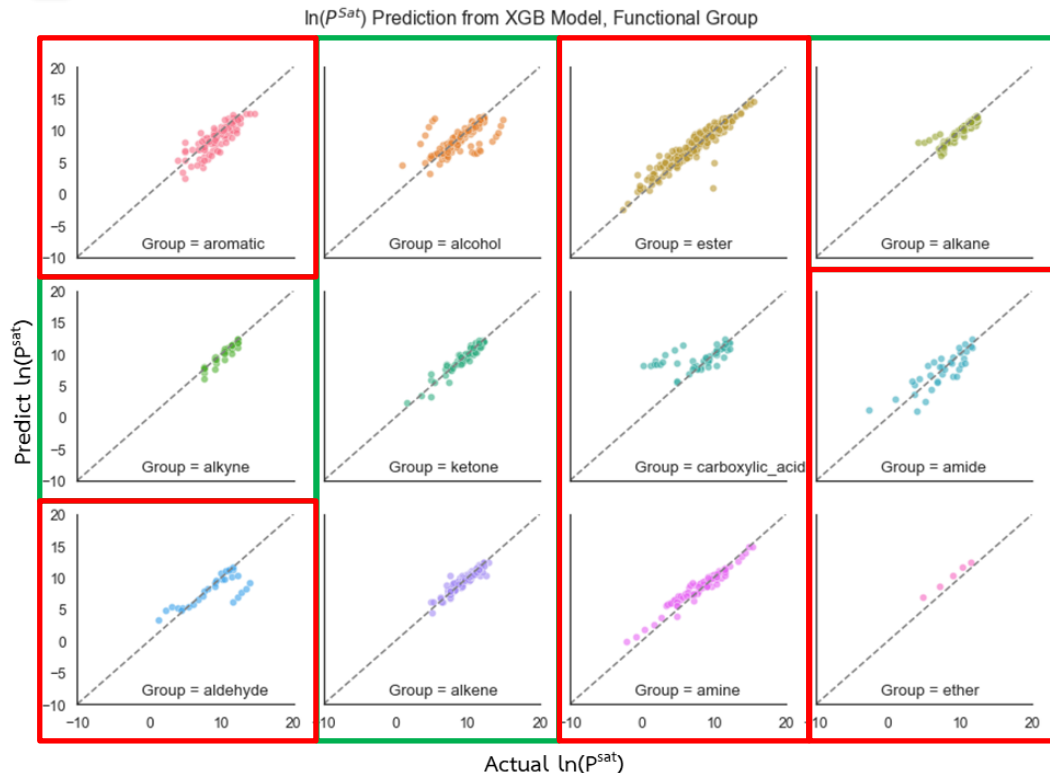
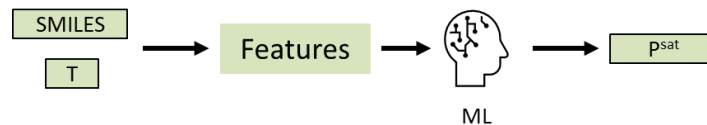


Table 8. Error metrics of each algorithm

Method	MAE	MAPE (%)	RMSE	R ²
DT	0.80	15.00	1.38	0.68
KNN	1.76	22.30	2.42	-1.24
RF	0.63	15.90	1.10	0.73
XGB	0.59	13.60	1.06	0.80

Result Evaluation



The functional group that this $\ln(p^{\text{sat}})$ prediction model can predict well is that the group that have condition with $R^2 \geq 0.5$ and $\text{MAPE} \leq 10\%$

Acceptable: Alcohol, Alkane, Alkyne, Alkene, Ketone

Unacceptable: Aromatic, Ester, Carboxylic Acid, Amide, Aldehyde, Amine, Ether

Fig 8. p^{sat} Prediction on CHON Scope Group by Functional Group

Result Calculation

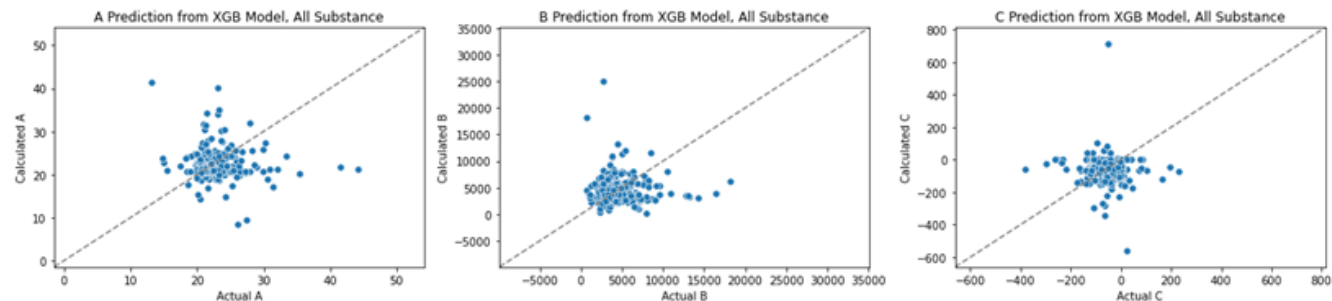
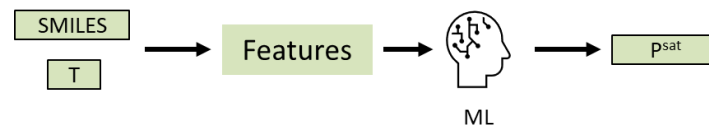
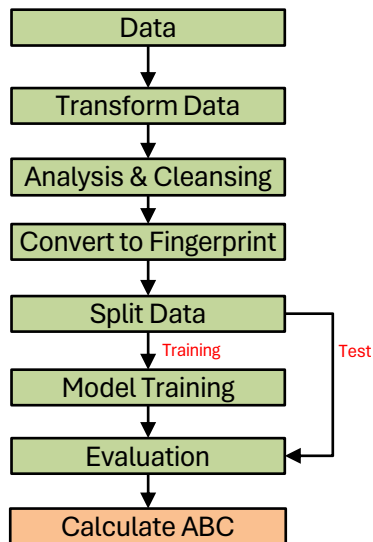


Fig 9. Graph Distribution of Antoine Coefficients

Calculate A, B, C for Predicted $\ln(P^{\text{sat}})$ from the best model

Result Calculation

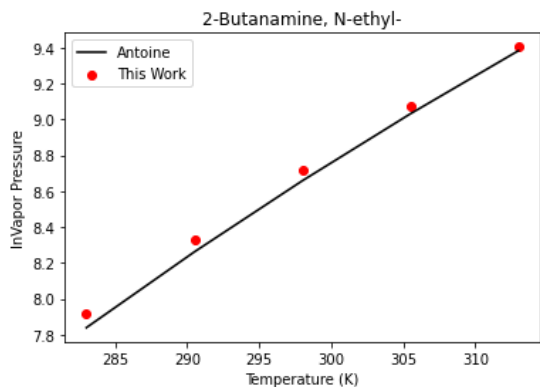
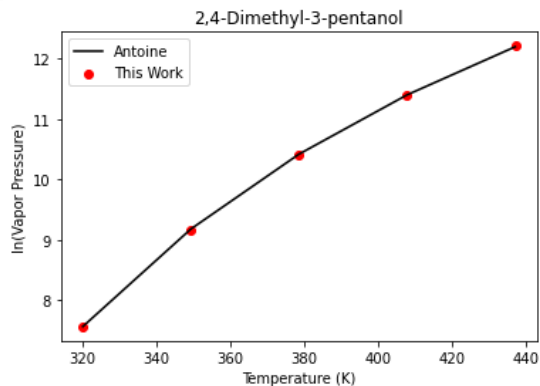


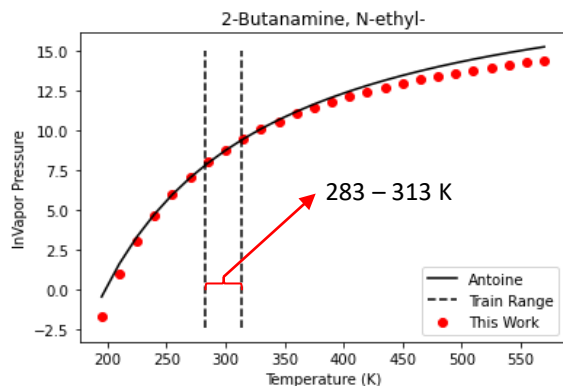
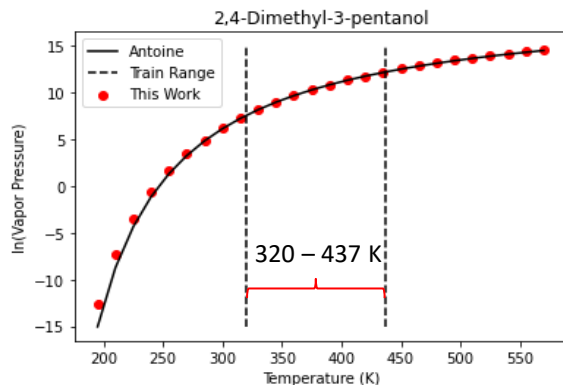
Table 8. Antoine Coefficients in this work and Antoine

SMILES	Reference			Calculation			Train Temp Range (K)
	A	B	C	A	B	C	
2,4-Dimethyl-3-pentanol	19.85	2370.74	-127.05	20.35	2615.59	-115.52	320 – 440
2-Butanamine, N-ethyl-	21.24	3083.56	-52.86	18.67	2006.16	-96.46	283 – 313

Antoine Coefficients that yield similar results to the reference source, even if slightly different, can still be used reliably.

Result

Calculation



When comparing temperatures at different range, it is observed that greater temperature differences range lead to more accurate predictions compared to the reference, especially when considering the lowest and highest temperatures from all references for all substances.

Table 9. Antoine Coefficients in this work and Antoine

SMILES	Reference			Calculation		
	A	B	C	A	B	C
2,4-Dimethyl-3-pentanol	19.85	2370.74	-127.05	20.35	2615.59	-115.52
2-Butanamine, N-ethyl-	21.24	3083.56	-52.86	18.67	2006.16	-96.46

Note: Temperature Range in Dataset (195 - 574 K)

Conclusion

1. Use Morgan Fingerprint to resolve Previous Work Problem “Similar Structures, get same Features”
2. Use Morgan Fingerprint to establish Properties Prediction Model (Use for Organic Compound C, H, O and N) : T_b - CHON Scope with MAPE = 5.94% and $R^2 = 0.66$
: $\ln(P^{sat})$ - CHON Scope with MAPE = 13.6% and $R^2 = 0.80$
3. Learn about Machine Learning Algorithm, Cross Validation and Hyperparameter Tuning in each Algorithm and Neural Network Implementation for Improving Properties Prediction model

Suggestions

1. Increasing the number of bits in a molecular fingerprint can help separate substructures, but some cases cannot be separated regardless of bit addition.
2. Deep Learning for modeling is more complex than Machine Learning, making it challenging to predict properties.
3. While modeling is faster, finding the right model takes time, so planning information preparation is crucial.
4. To improve model accuracy, consider creating a model that separates specific boundaries, focusing on specific functional groups.



Q&A