

Practical Machine Learning for Organic Small Molecule Modeling

Machine Learning Modalities for Materials Science Workshop

16 May 2024

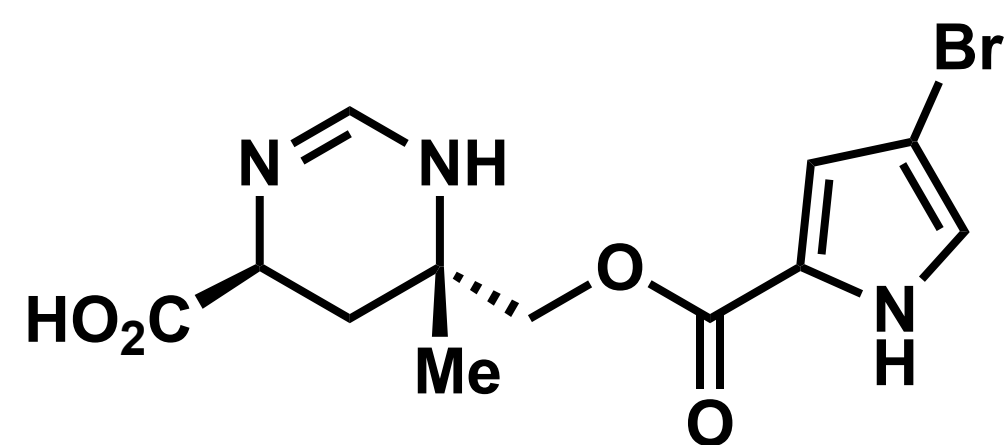
Emma King-Smith

What Do Synthetic Chemists Want?

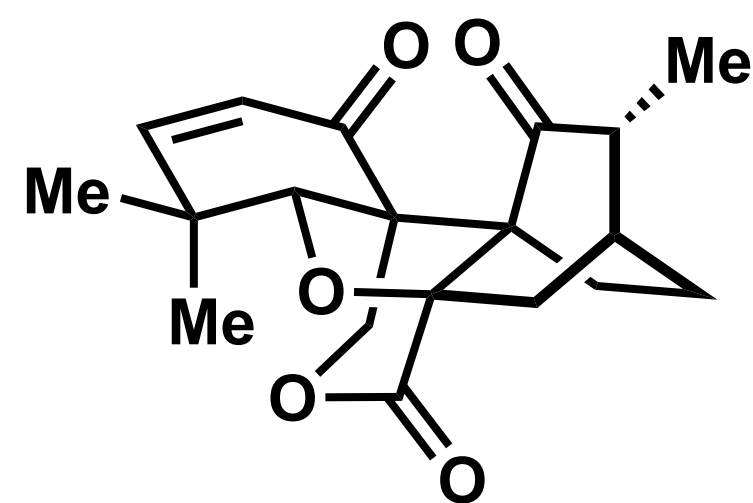
Make Molecules!

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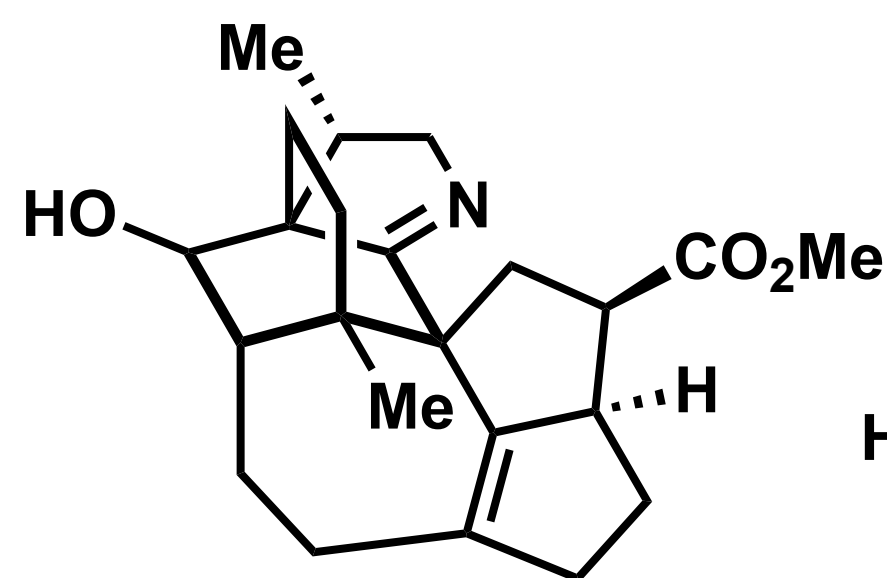
“Pretty” Molecules



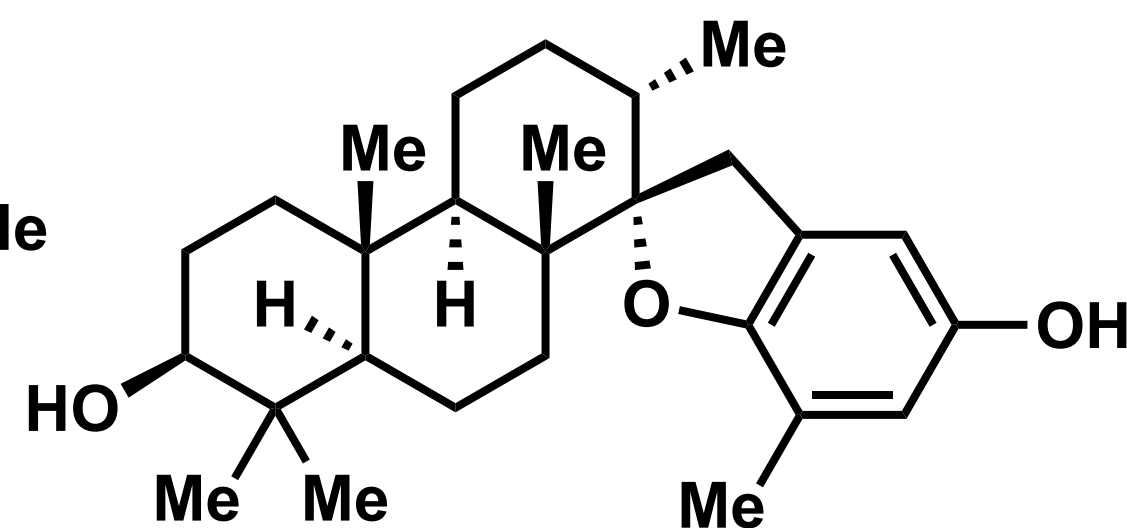
manzacidin C



maoecrystal V



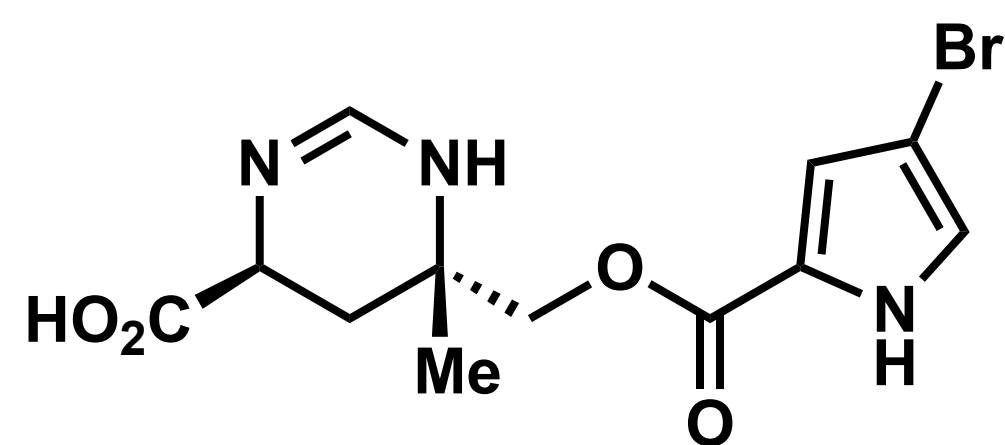
(-)-calyciphylline N



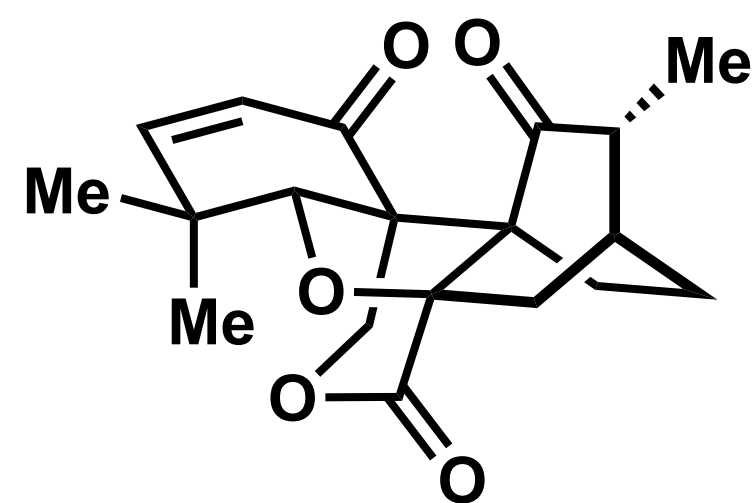
stypodiol

Make Molecules!

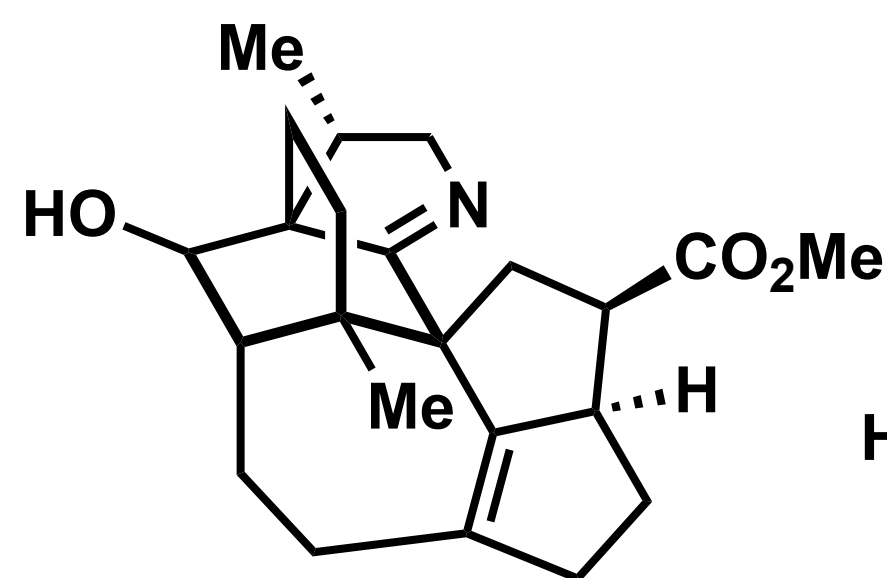
“Pretty” Molecules



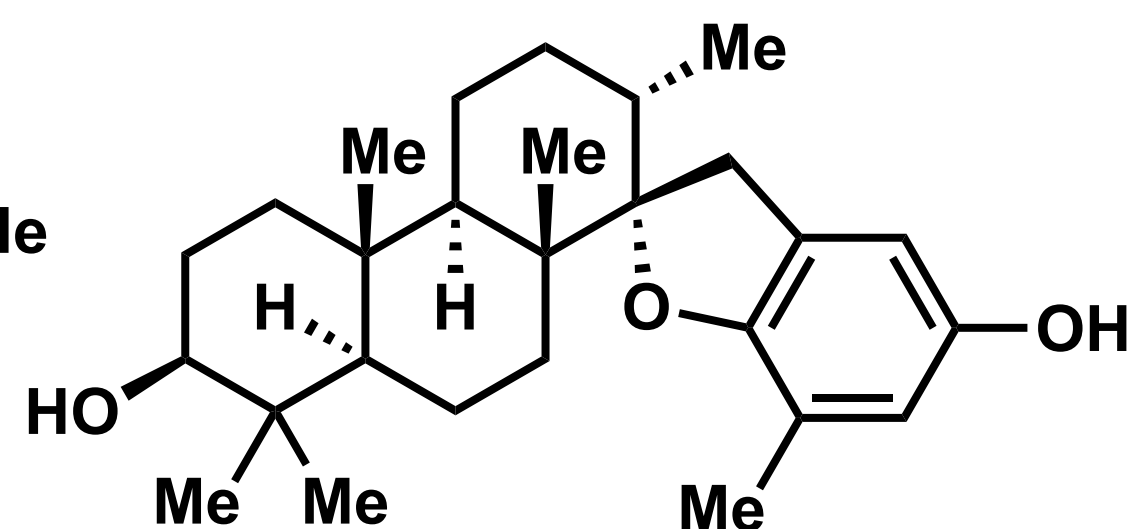
manzacidin C



maoecrystal V

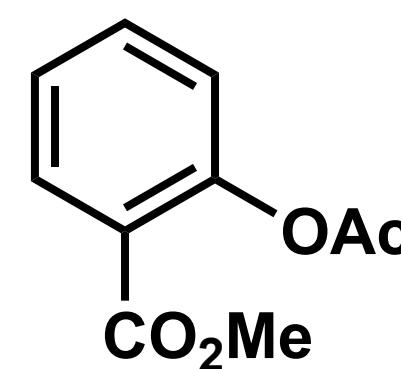


(-)-calyciphylline N

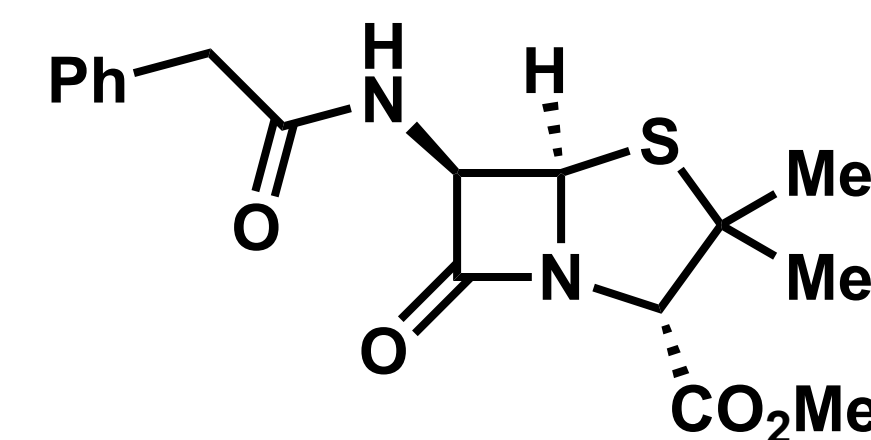


stypodiol

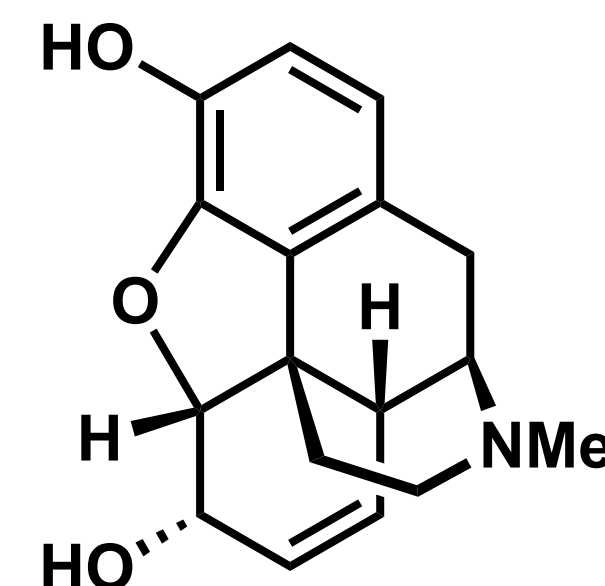
Pharmaceuticals



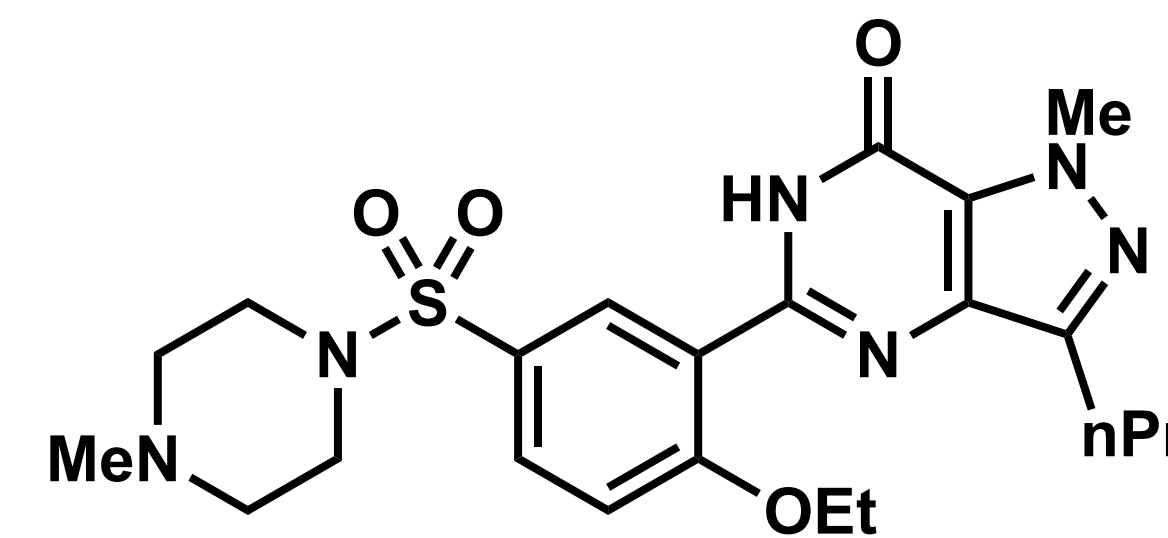
aspirin



penicillin G



morphine

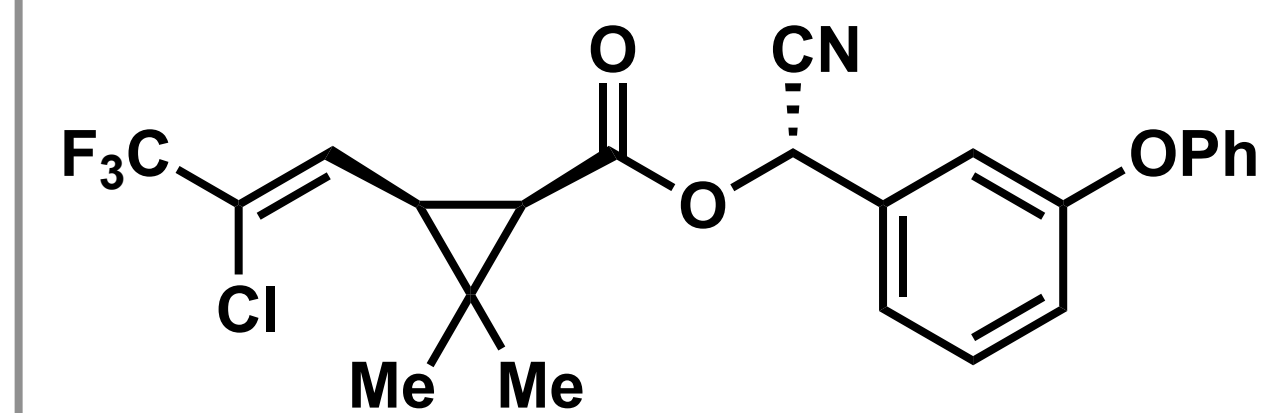


sildenafil

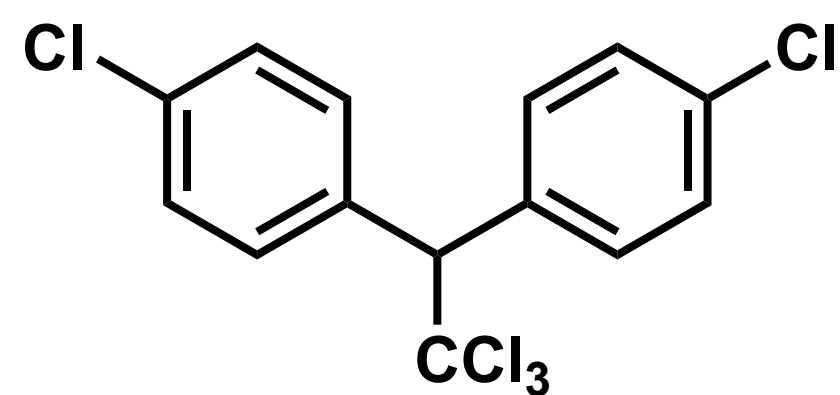
What Do Synthetic Chemists Want?

Make Molecules!

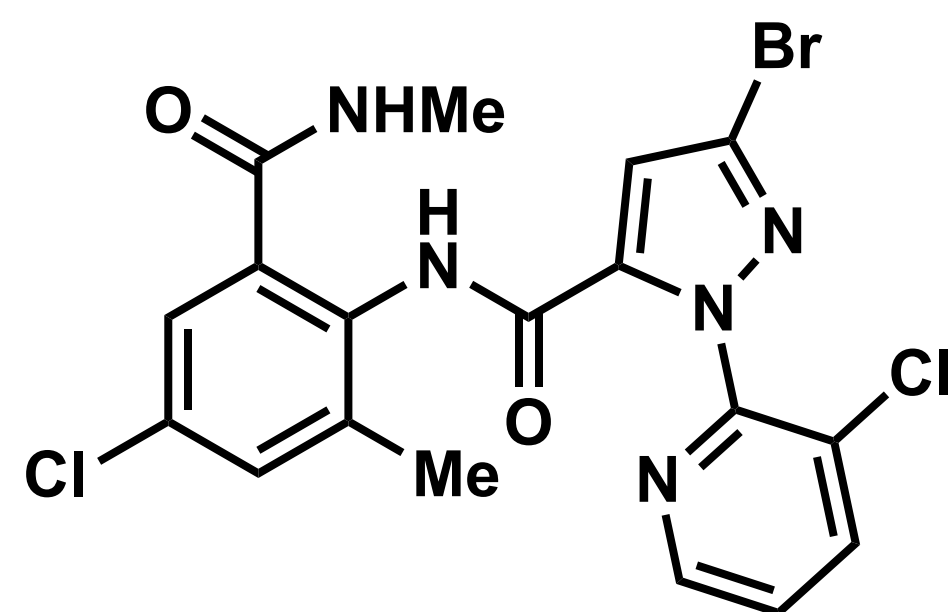
Agrochemicals



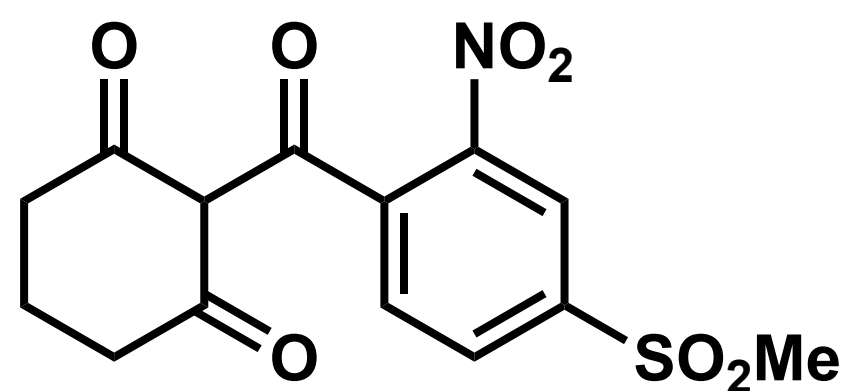
lambda-cyhalothrin



DDT



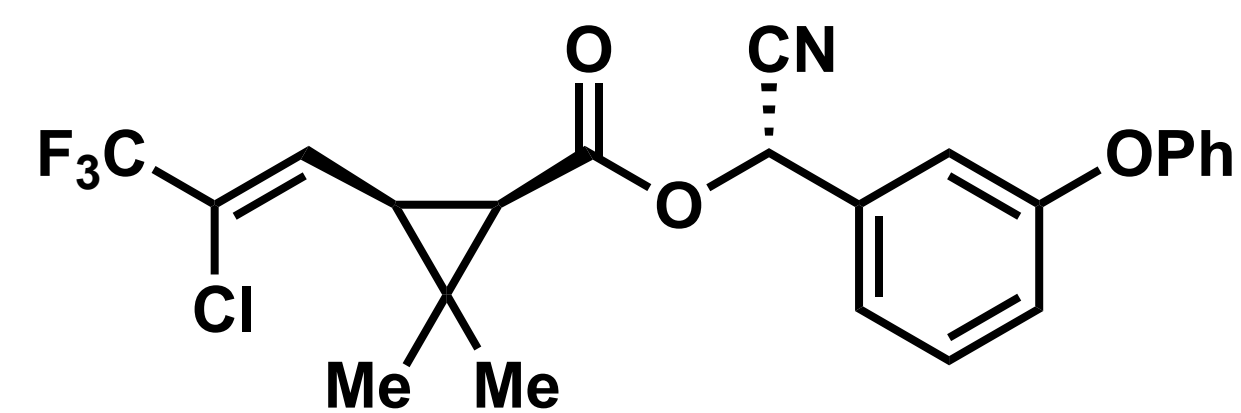
chlorantraniliprole



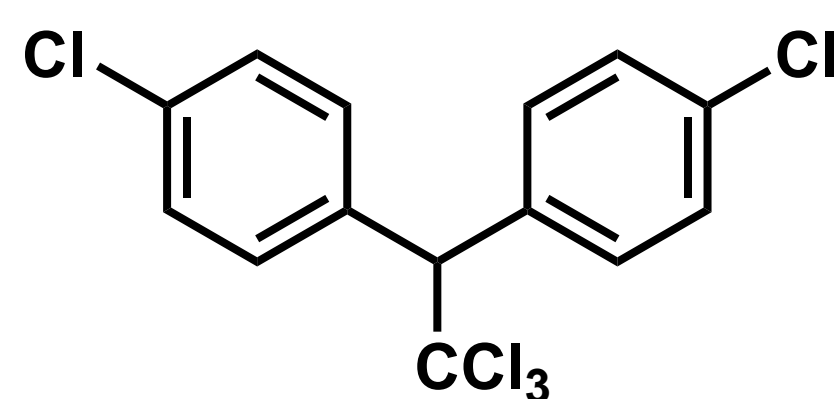
mesotrione

Make Molecules!

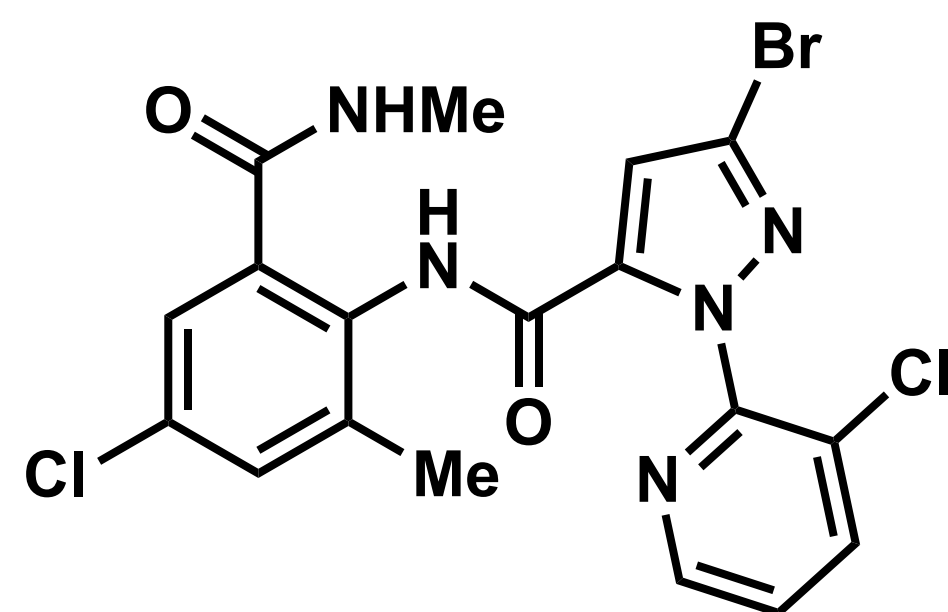
Agrochemicals



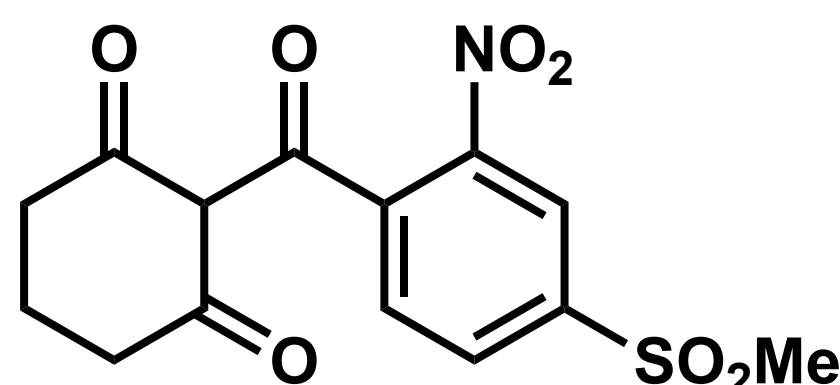
lambda-cyhalothrin



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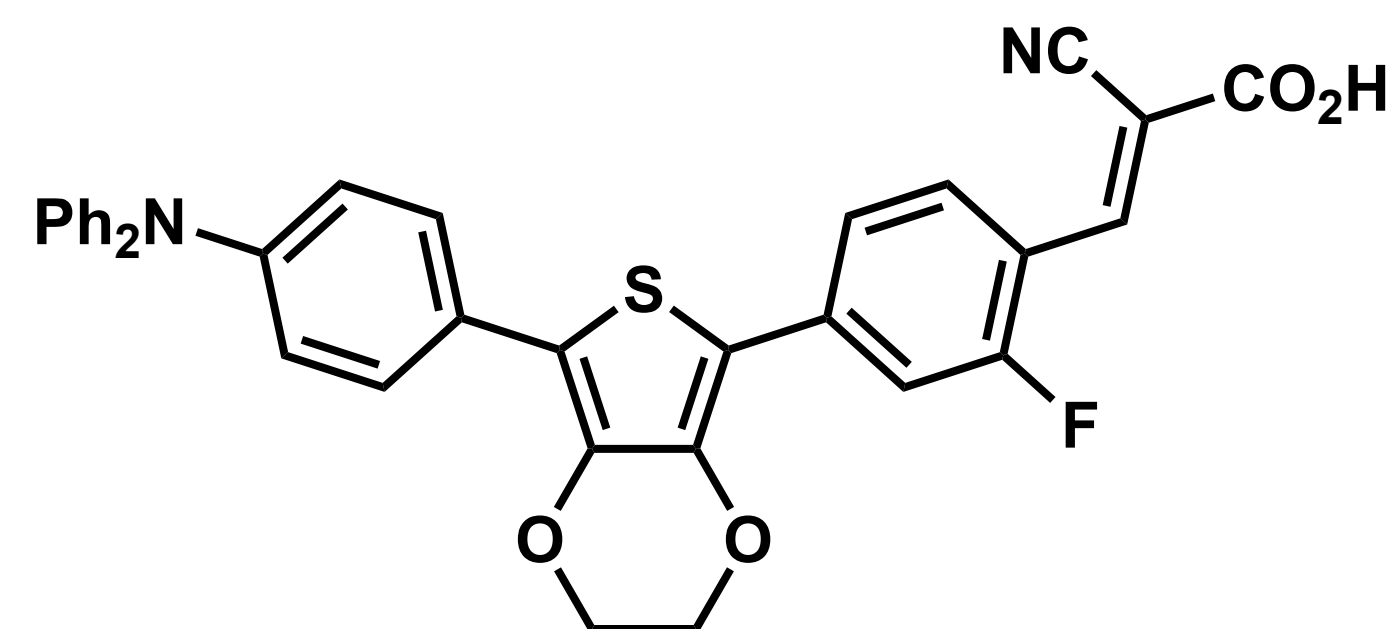


chlorantraniliprole

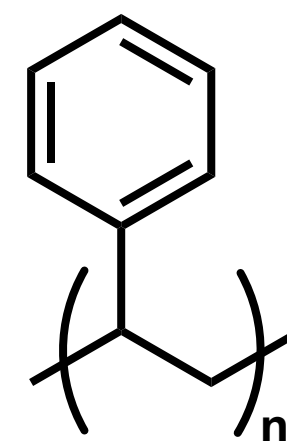


mesotrione

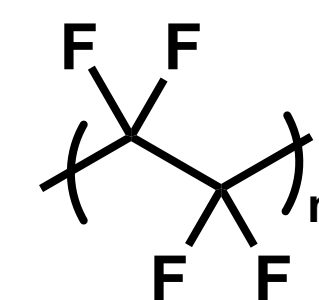
Organic Materials



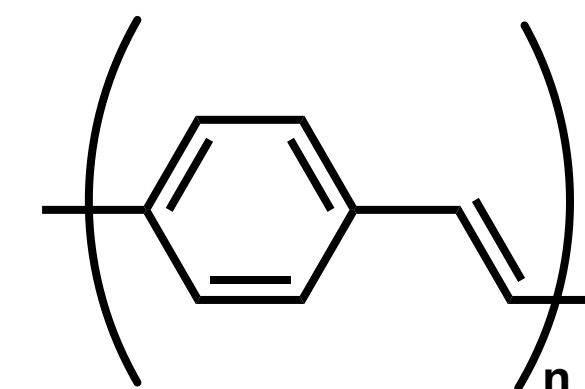
LBJ-F₀



polystyrene



Teflon



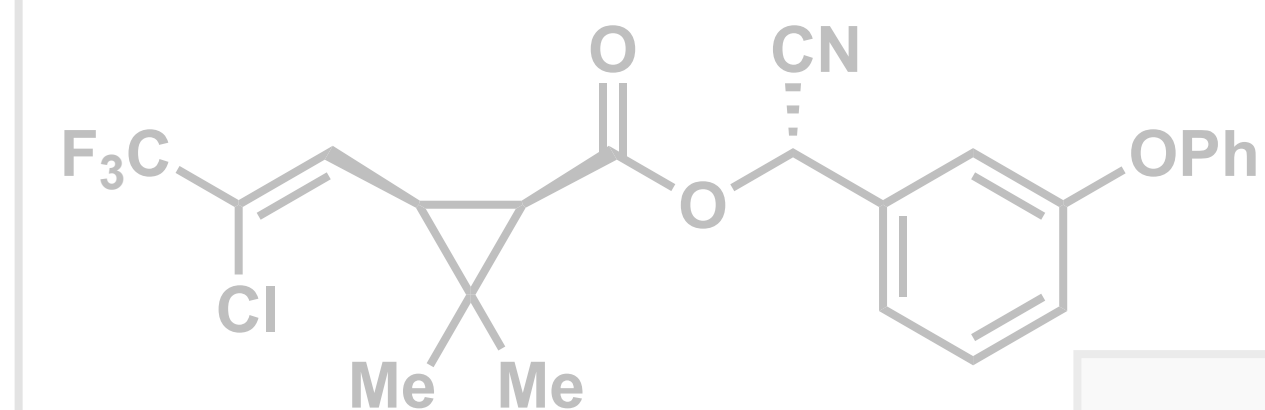
poly(*p*-phenylenevinylene)

What Do Synthetic Chemists Want?

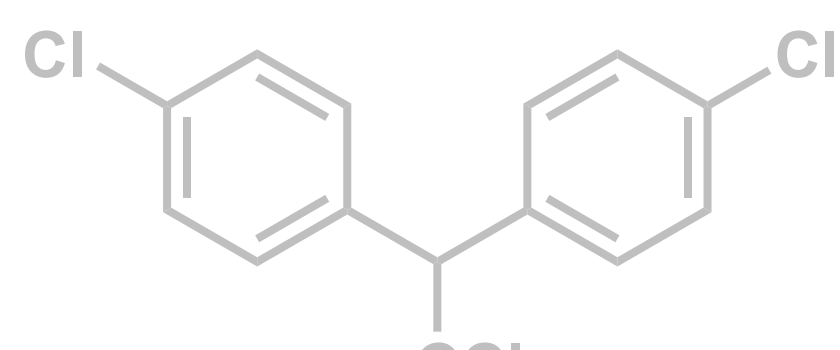
Make Molecules!

Agrochemicals

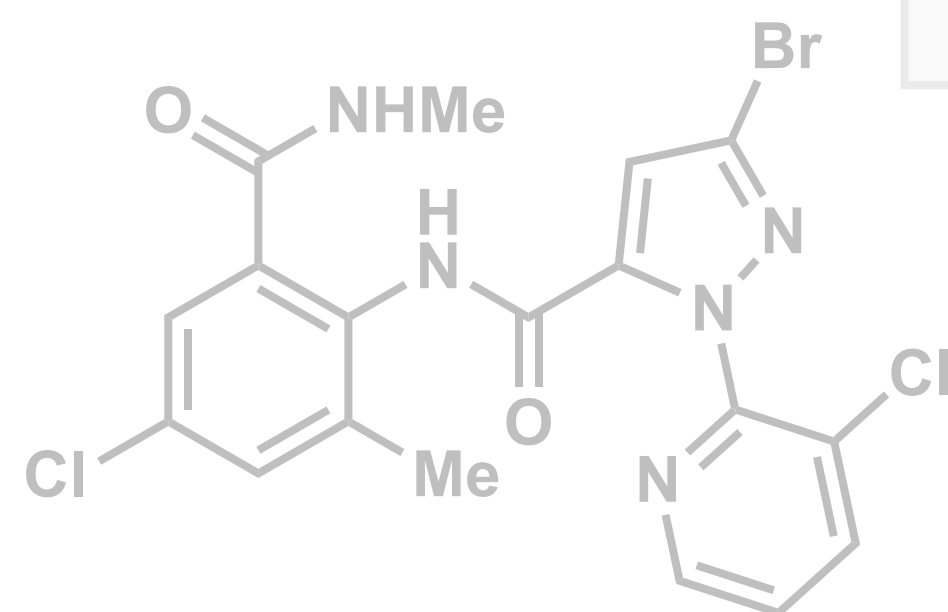
Organic Materials



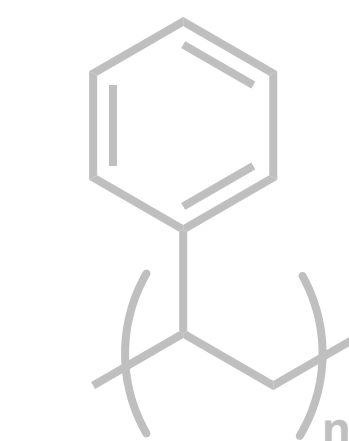
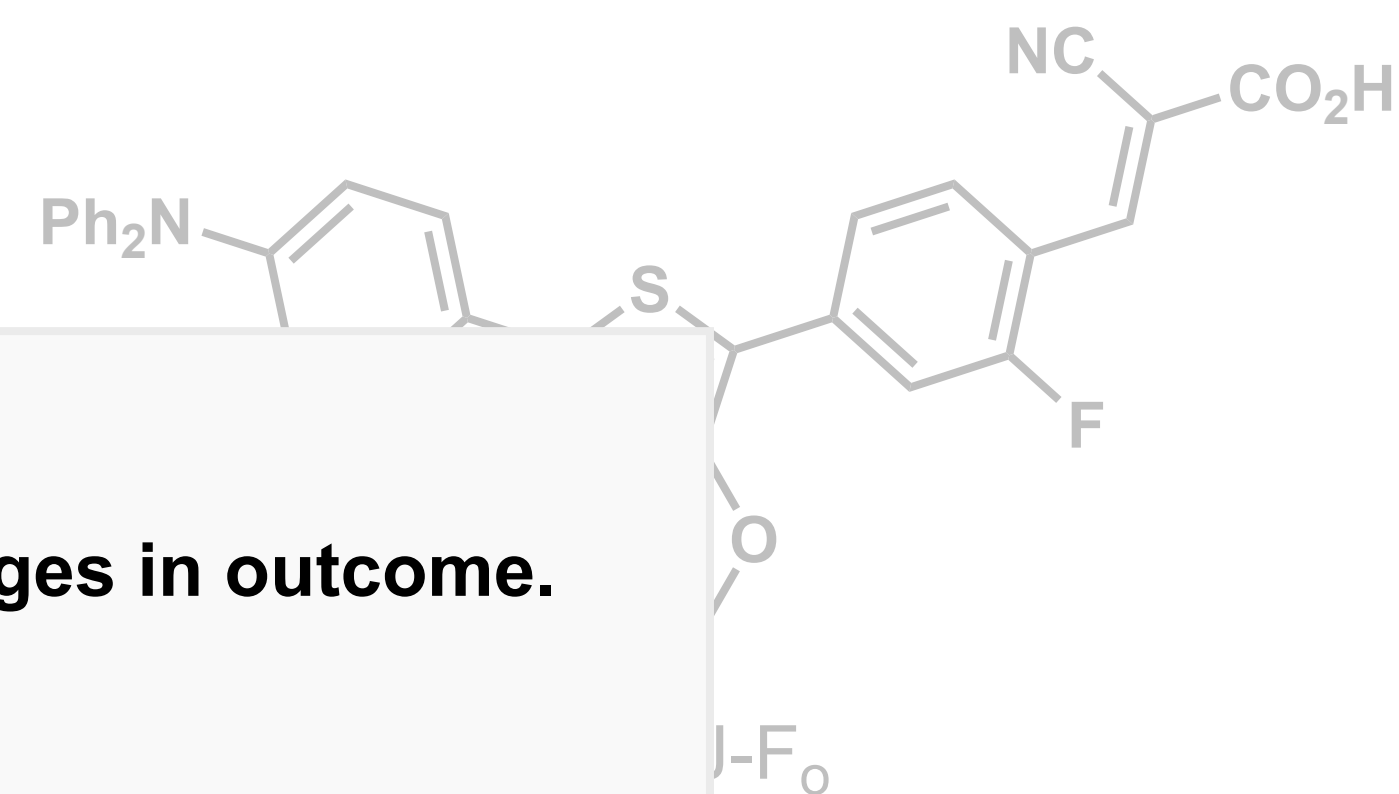
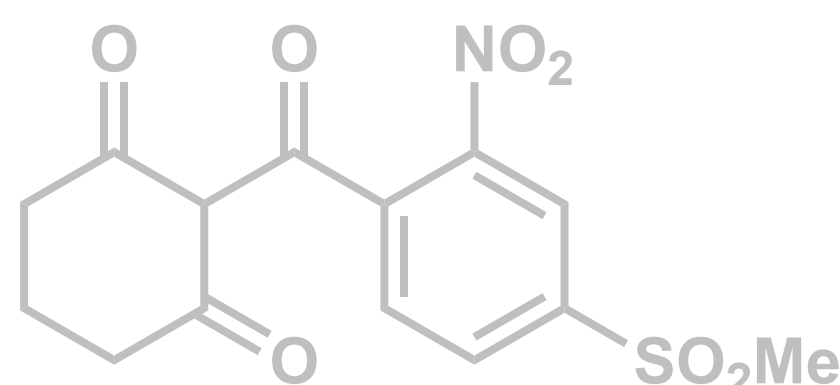
lambda-cyhalothrin



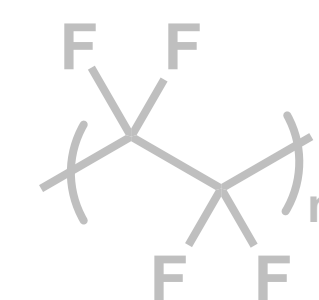
mesotrione



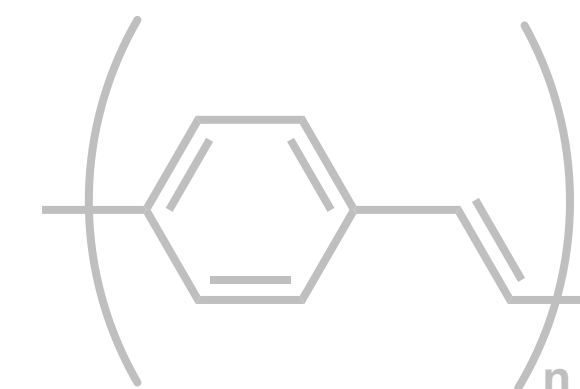
chlorantraniliprole



polystyrene



Teflon



poly(p-phenylenevinylene)

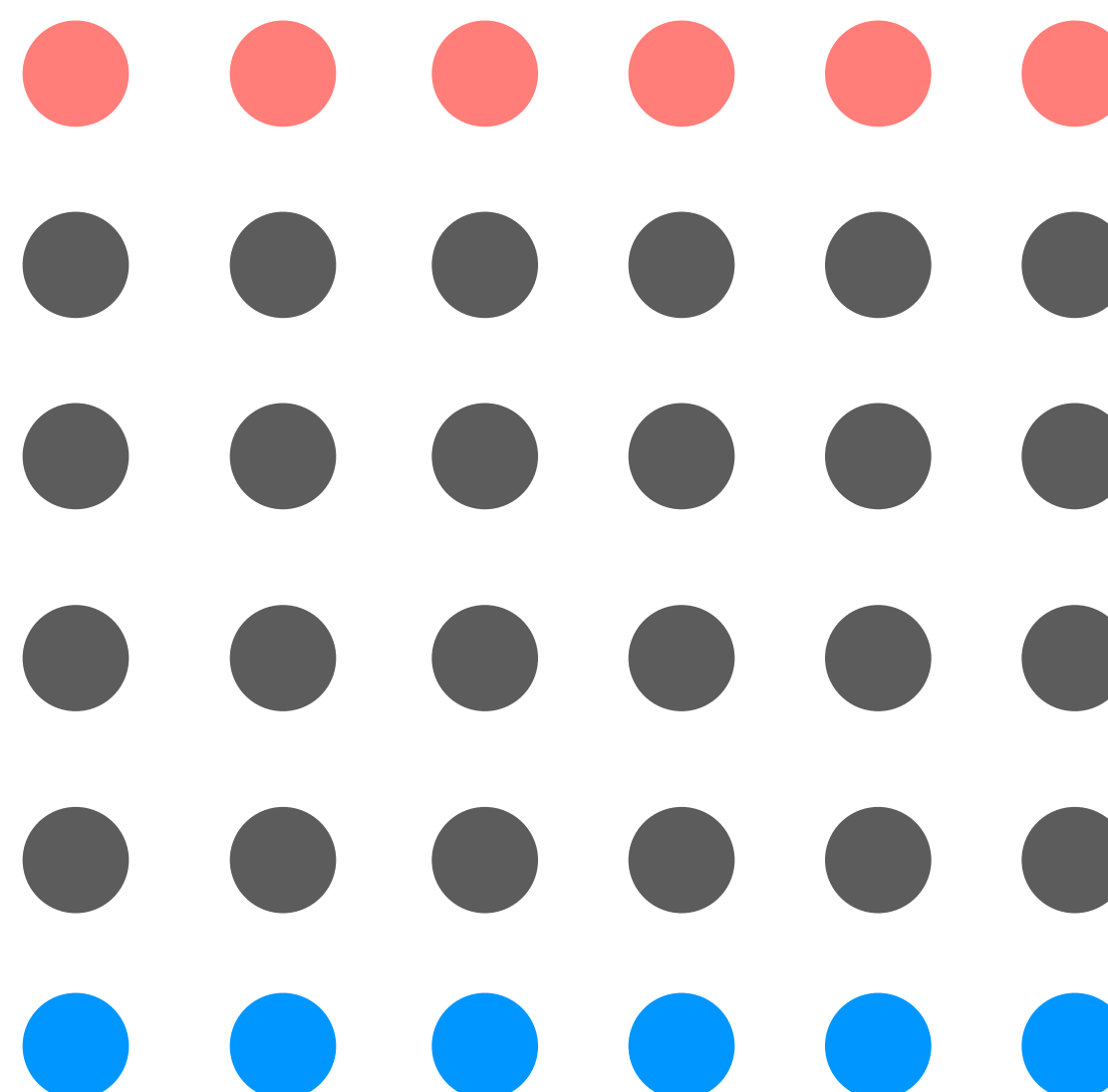
Small changes in structure ≠ Small changes in outcome.

Part I:

Transfer Learning to Unlock Chemical Predictions in Low Data
Regimes

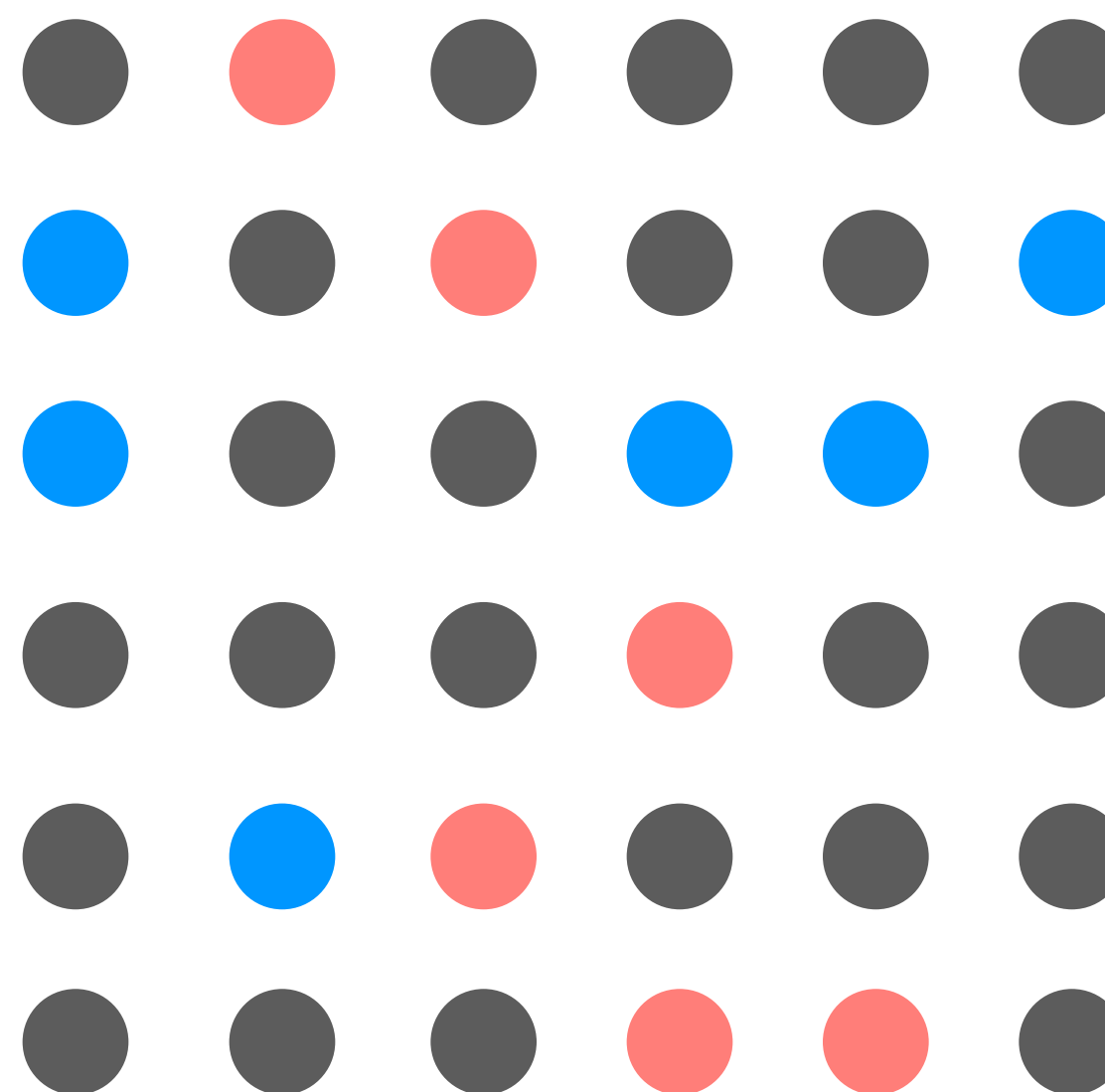
The Importance of Test Sets

● = molecule 1 ● = molecule 2 ● = molecule 3



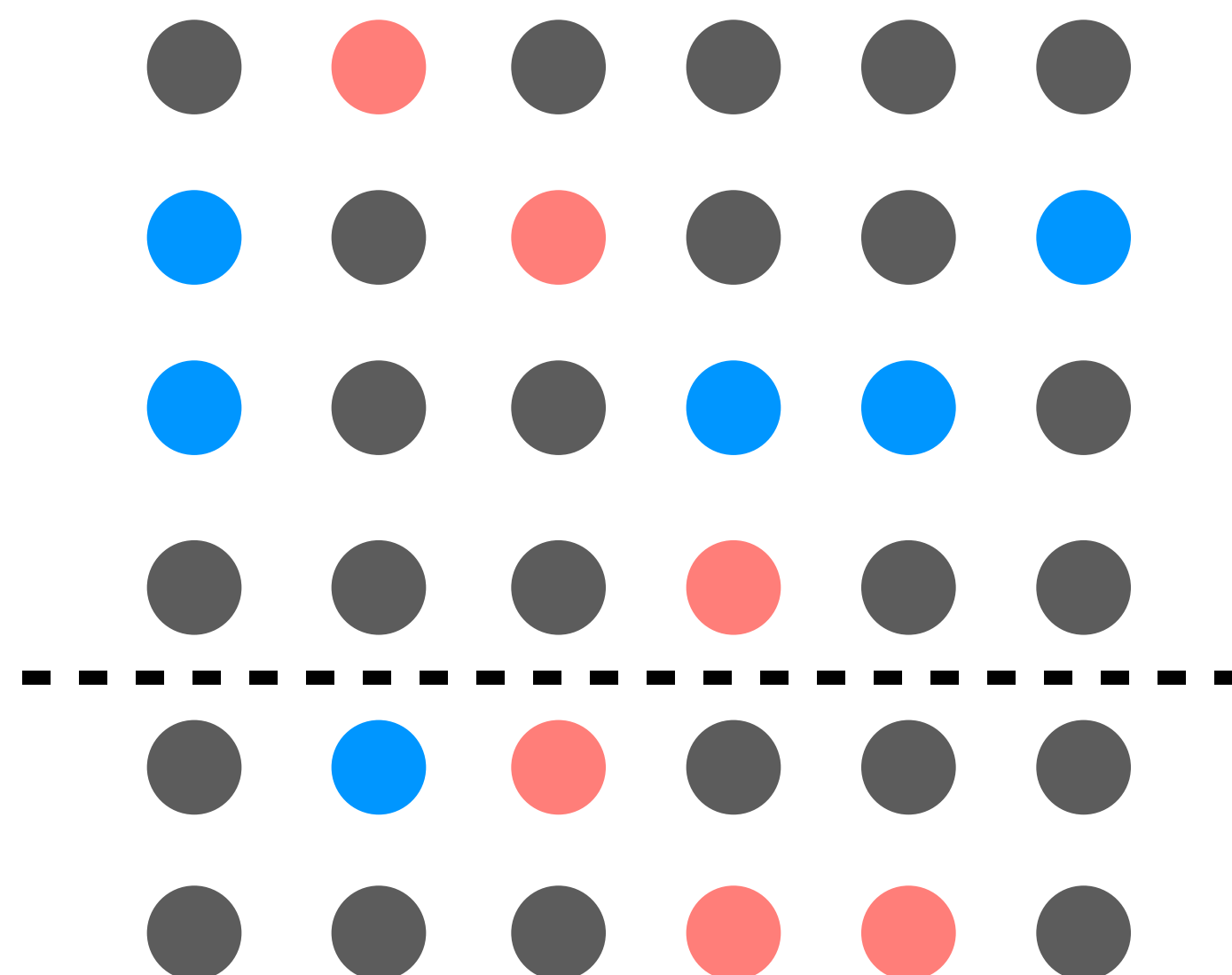
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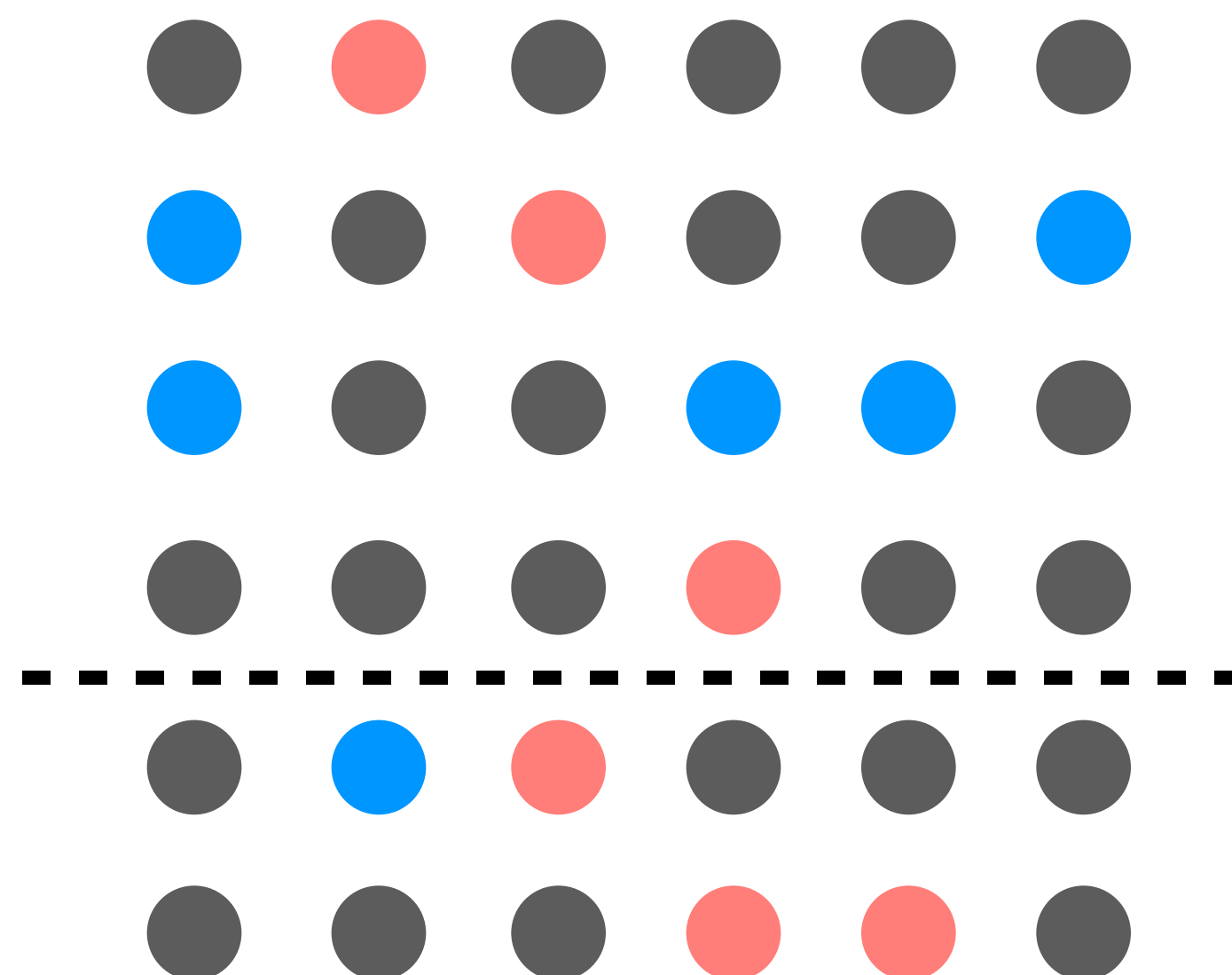
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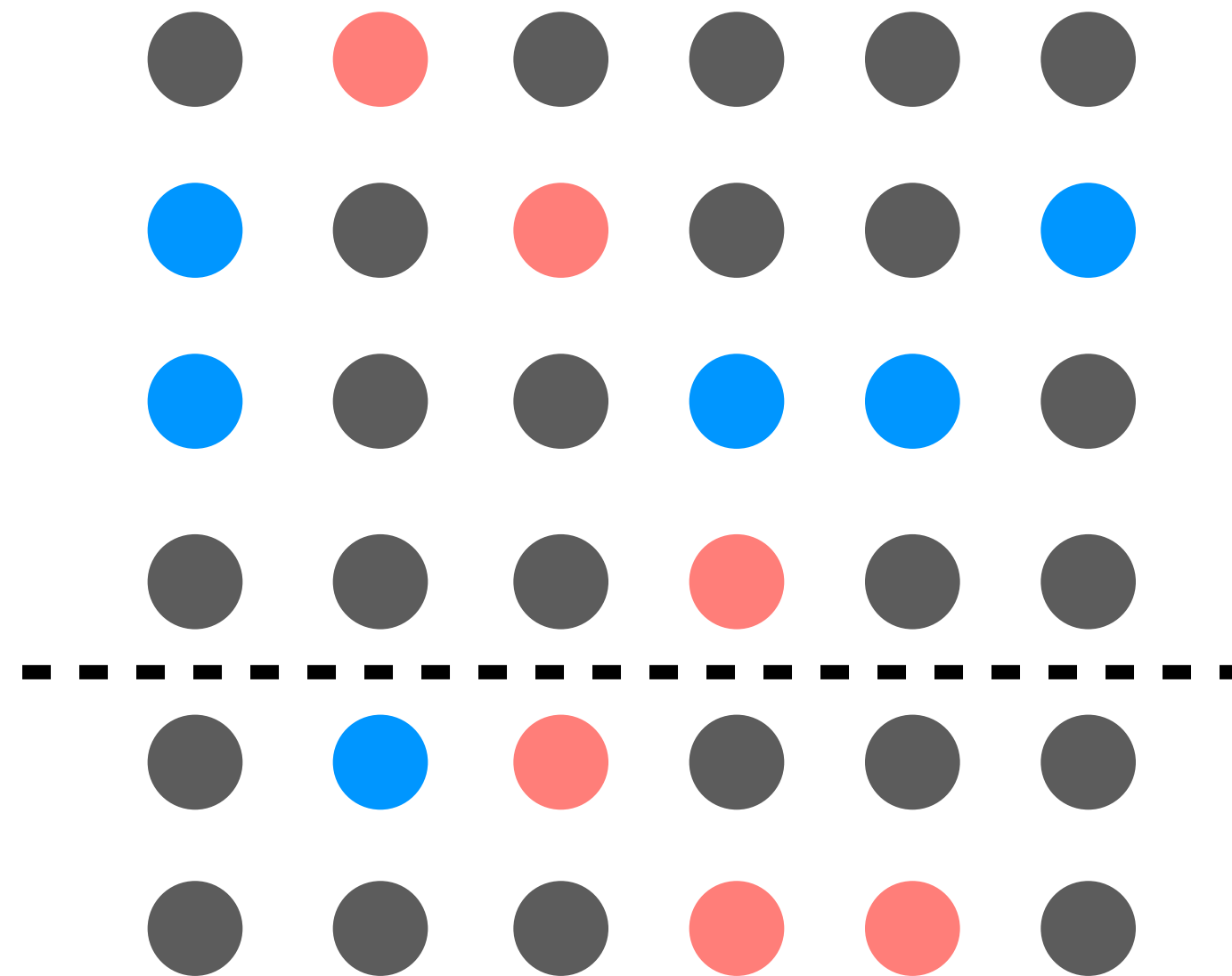
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simple

The Importance of Test Sets

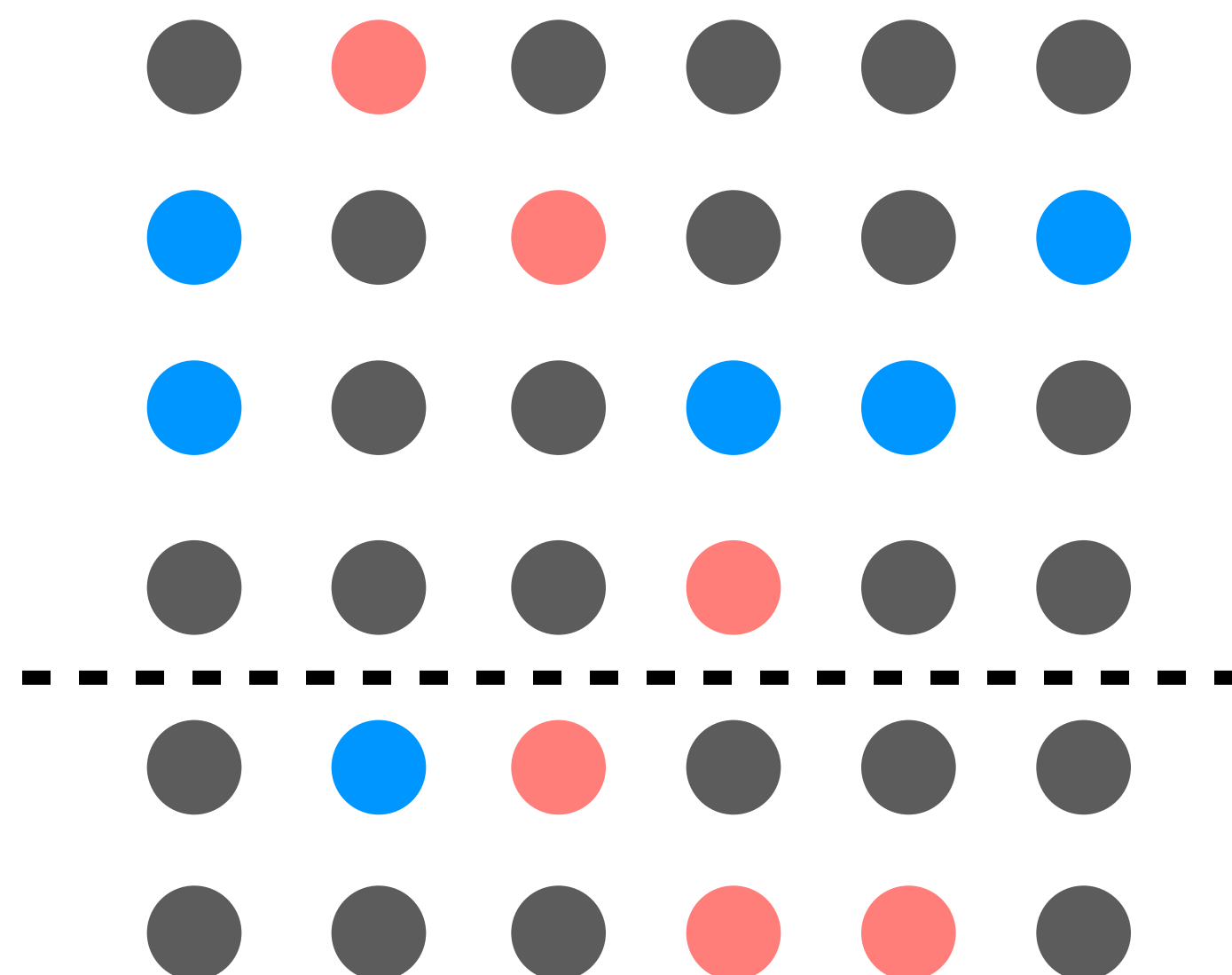
● = molecule 1 ● = molecule 2 ● = molecule 3



simple

easy to implement

● = molecule 1 ● = molecule 2 ● = molecule 3



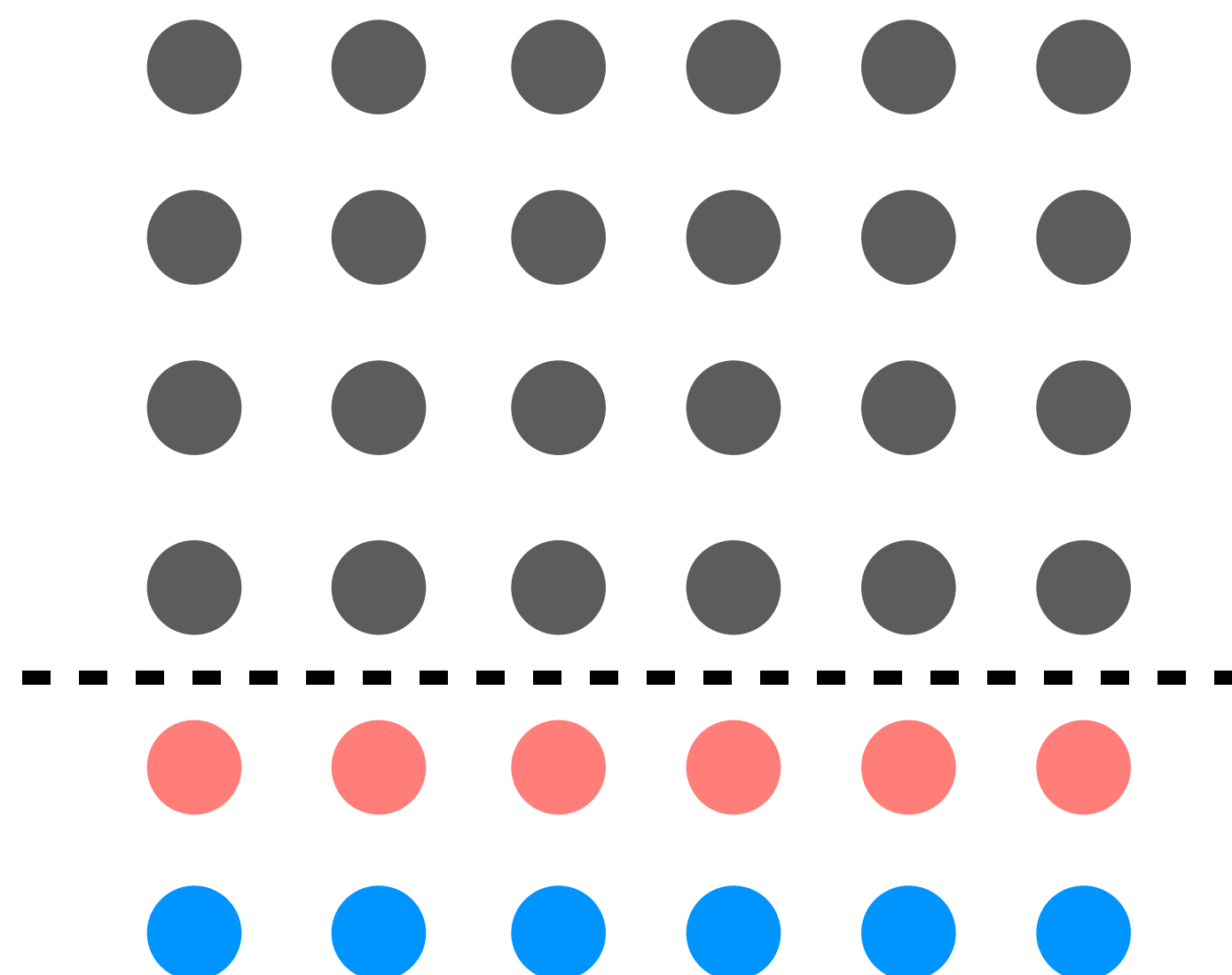
simple

easy to implement

can be used with any dataset

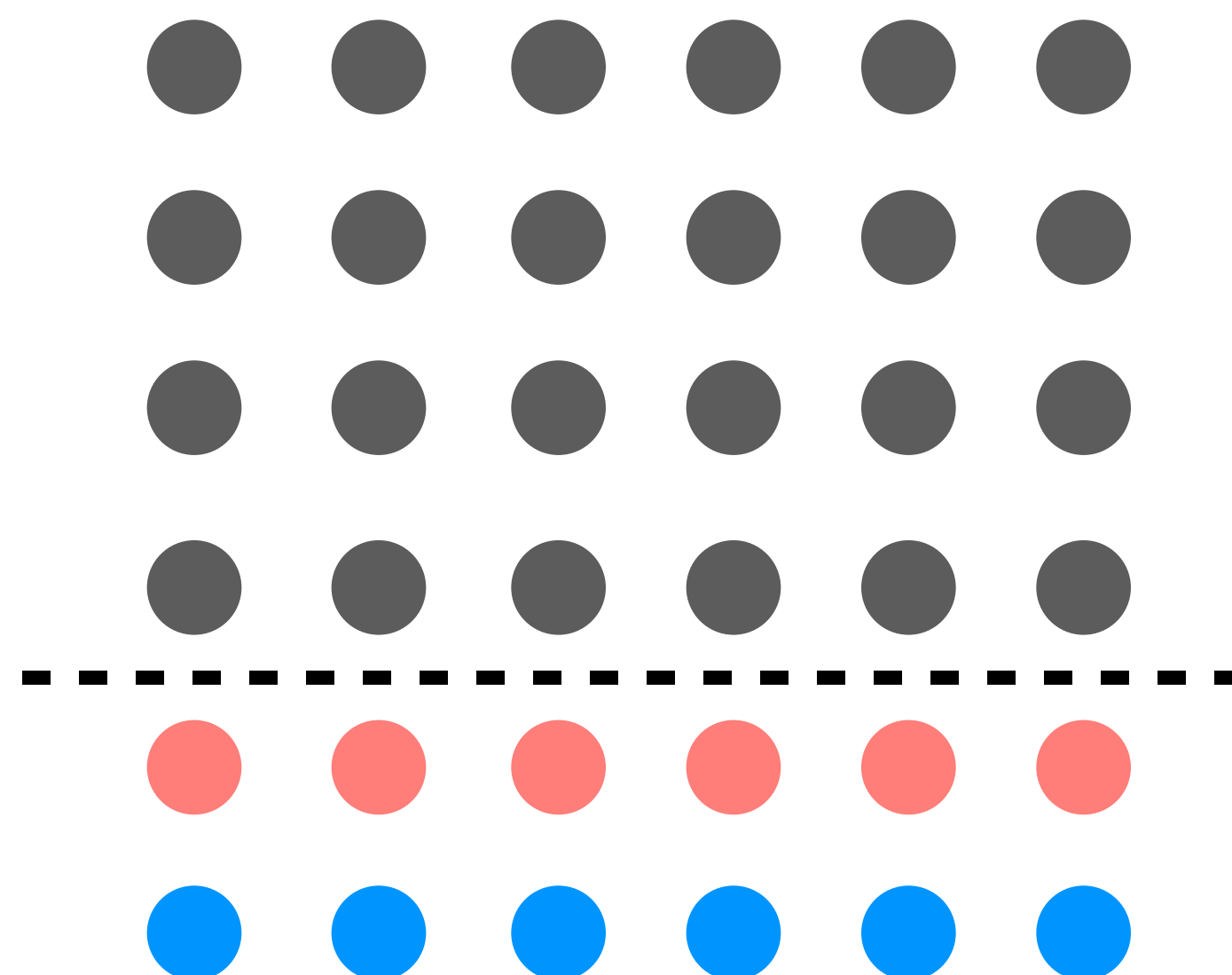
The Importance of Test Sets

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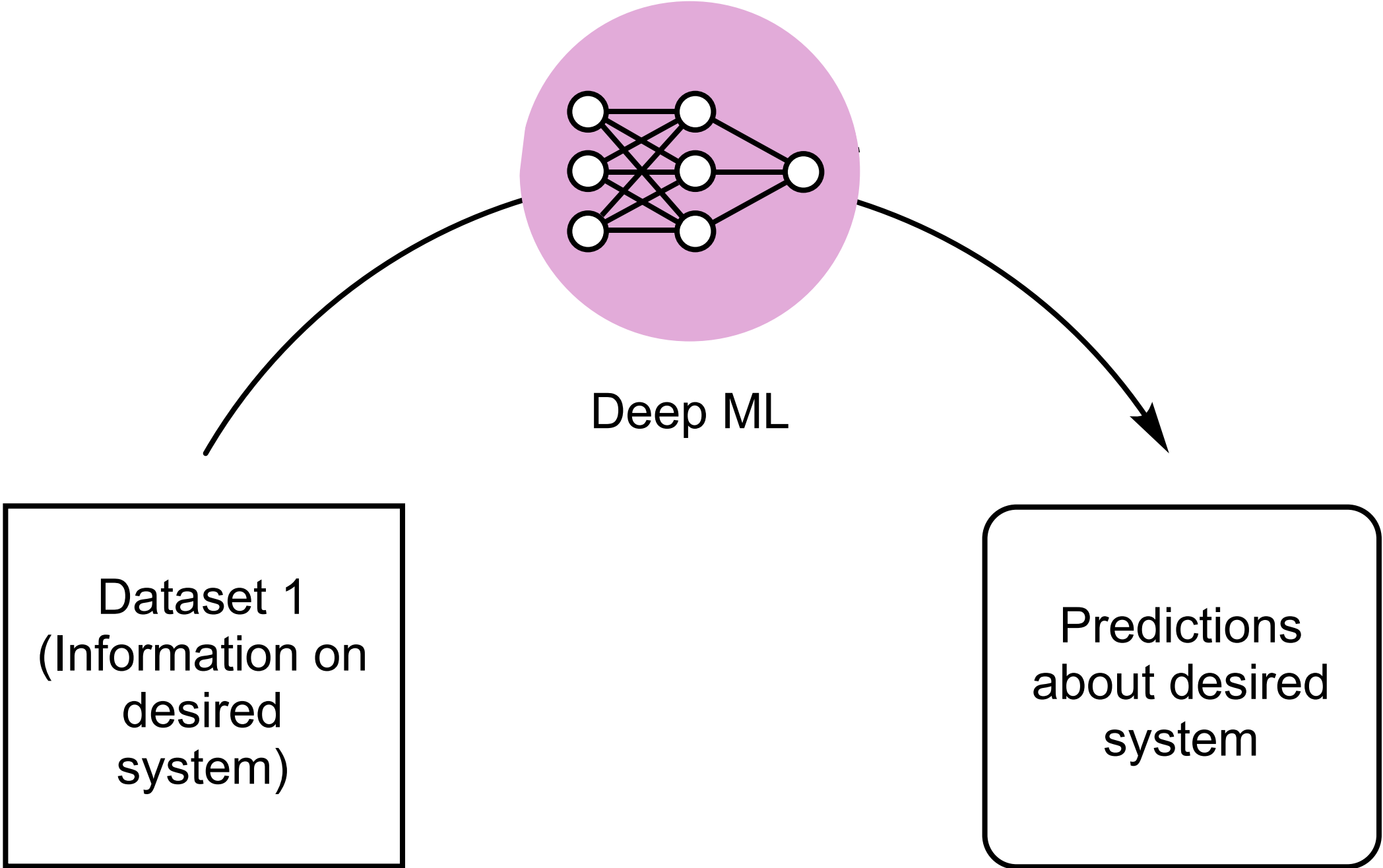
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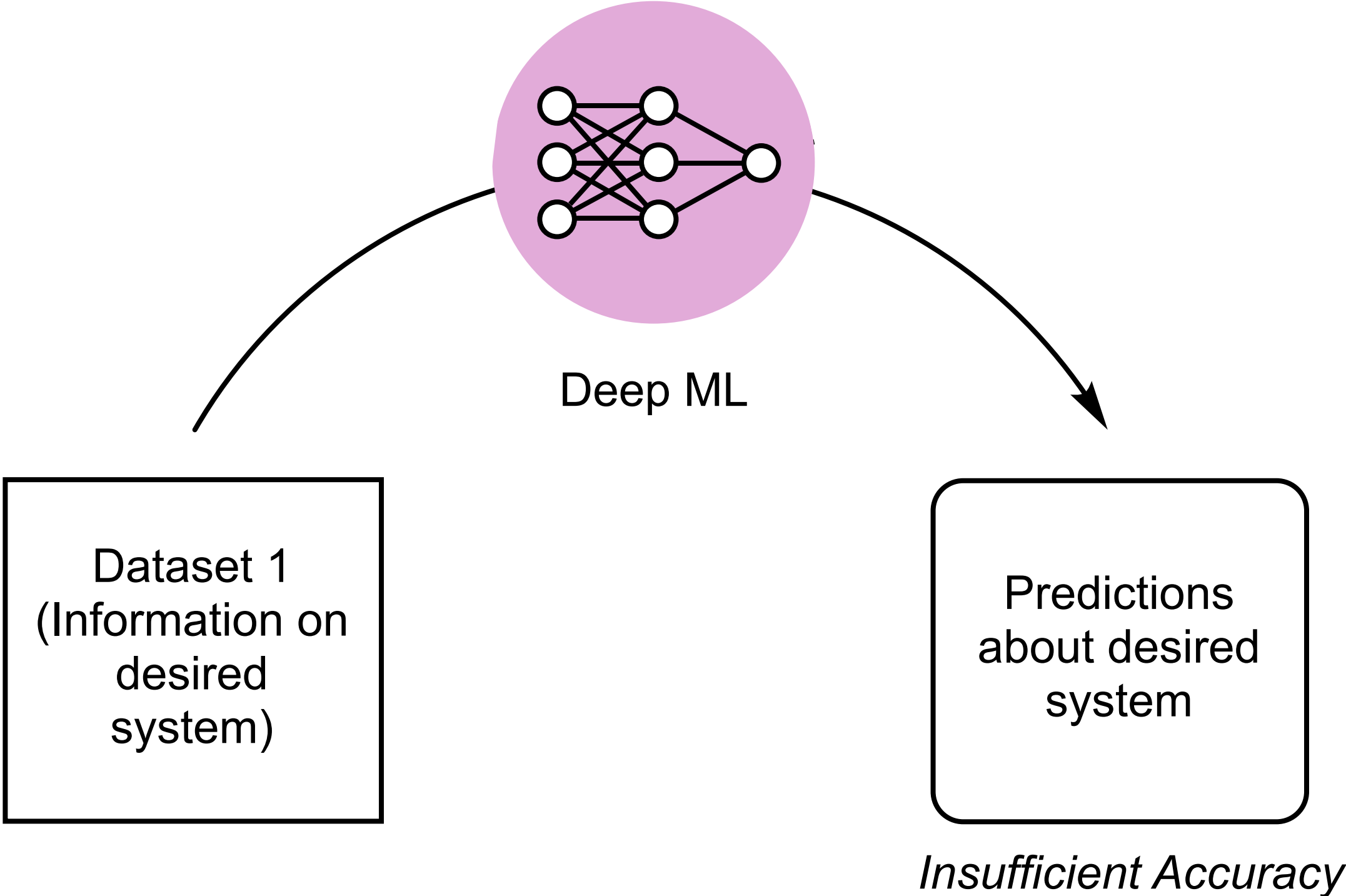


more challenging

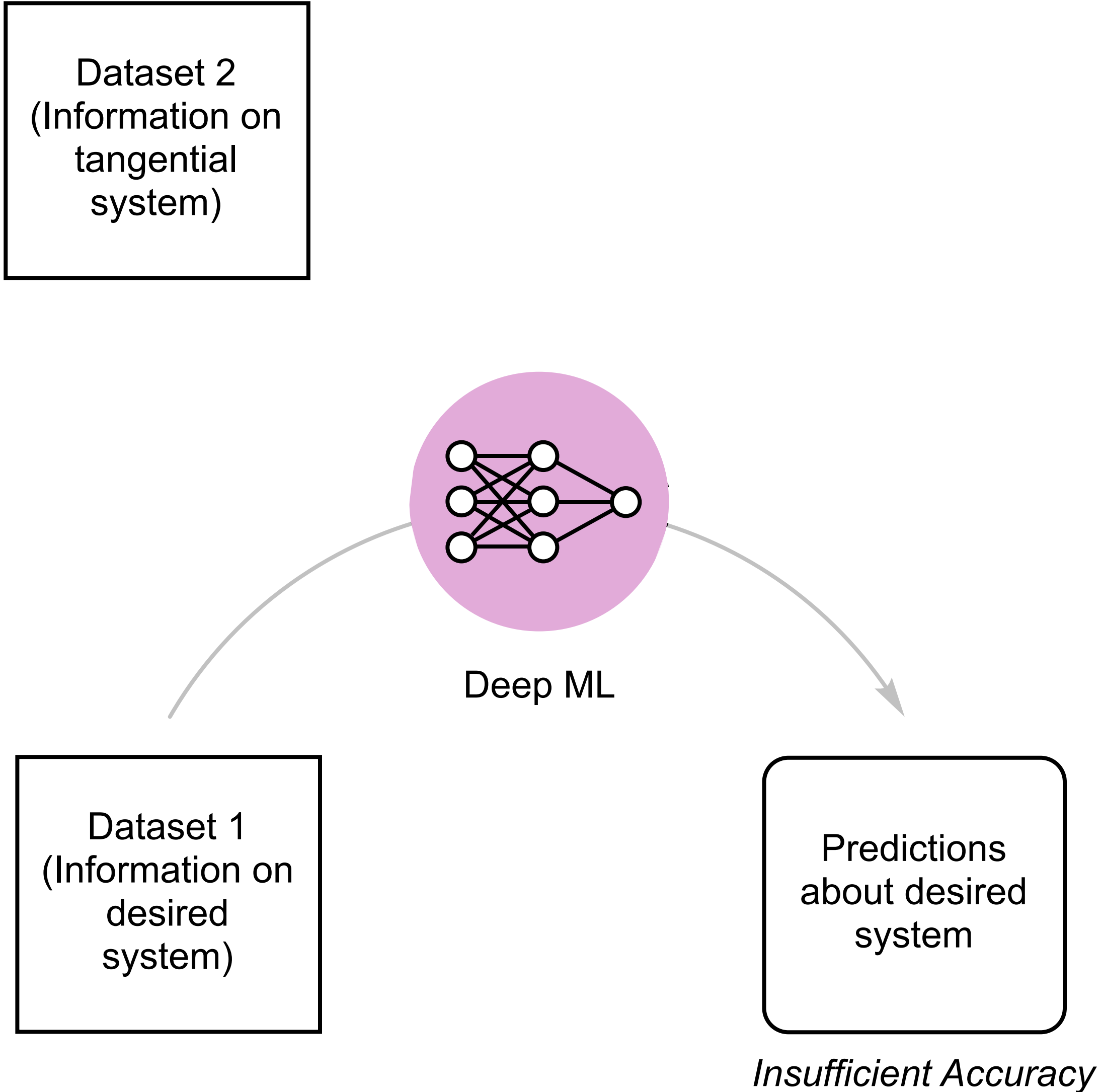
What Is Transfer Learning?



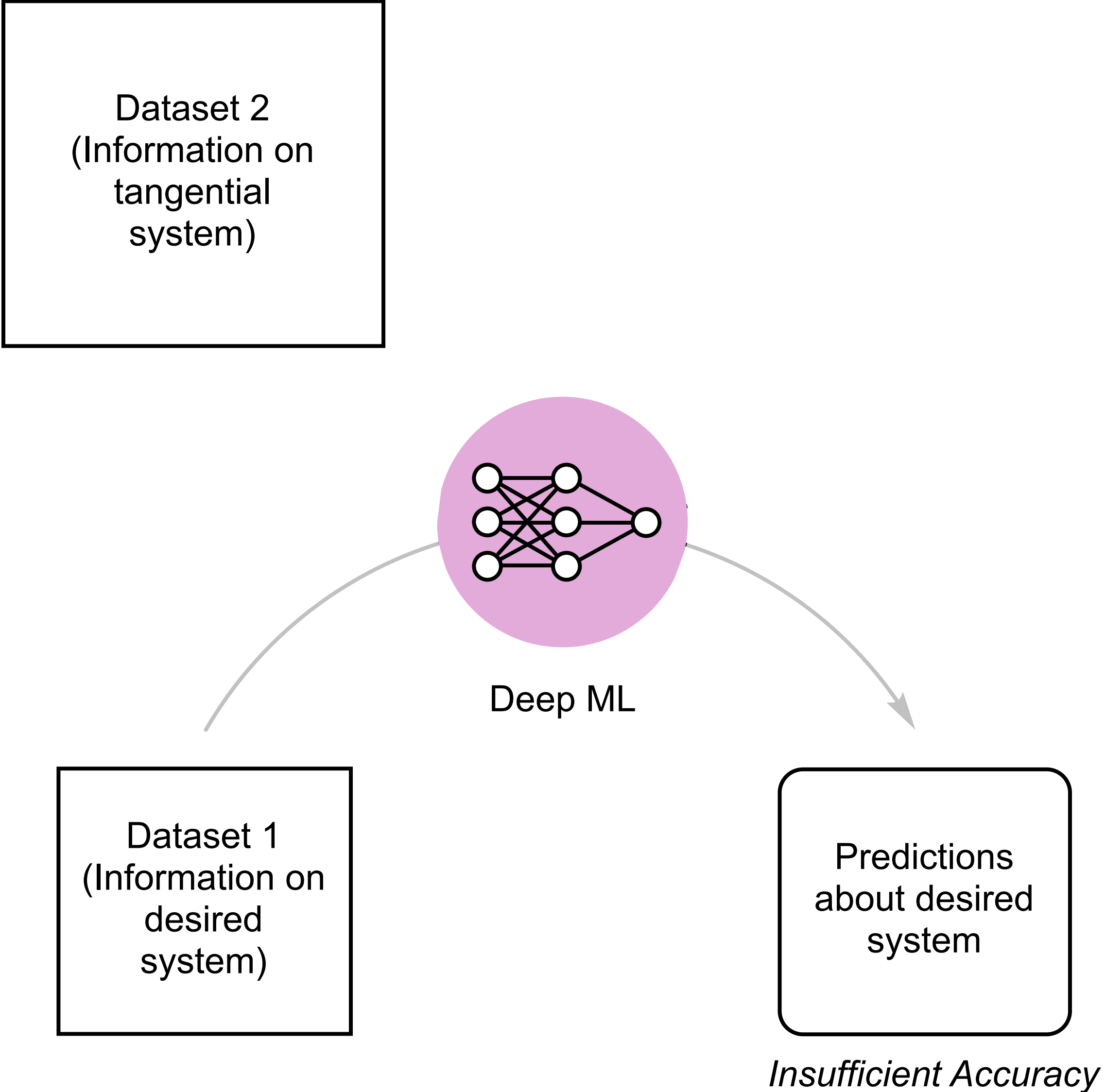
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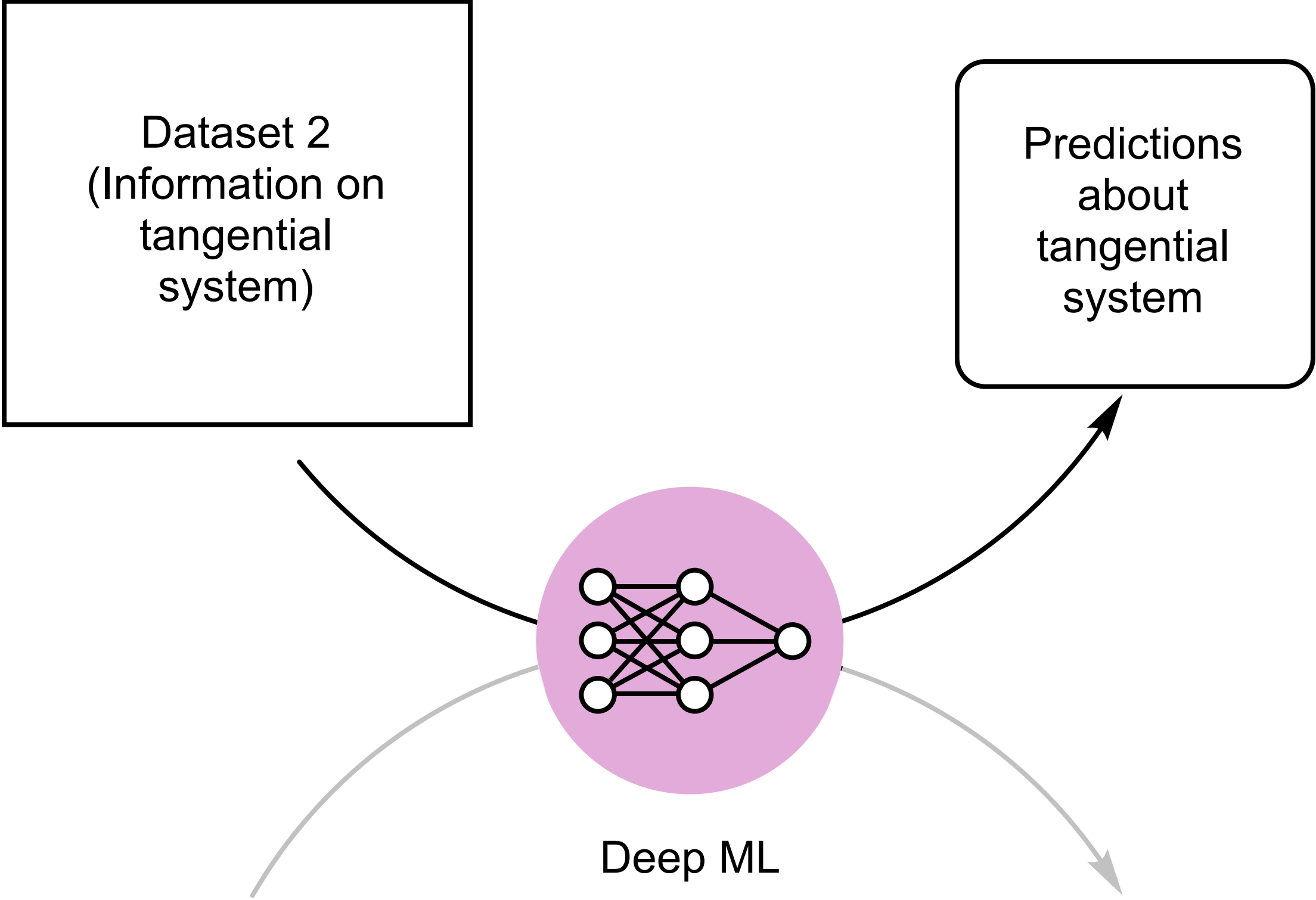
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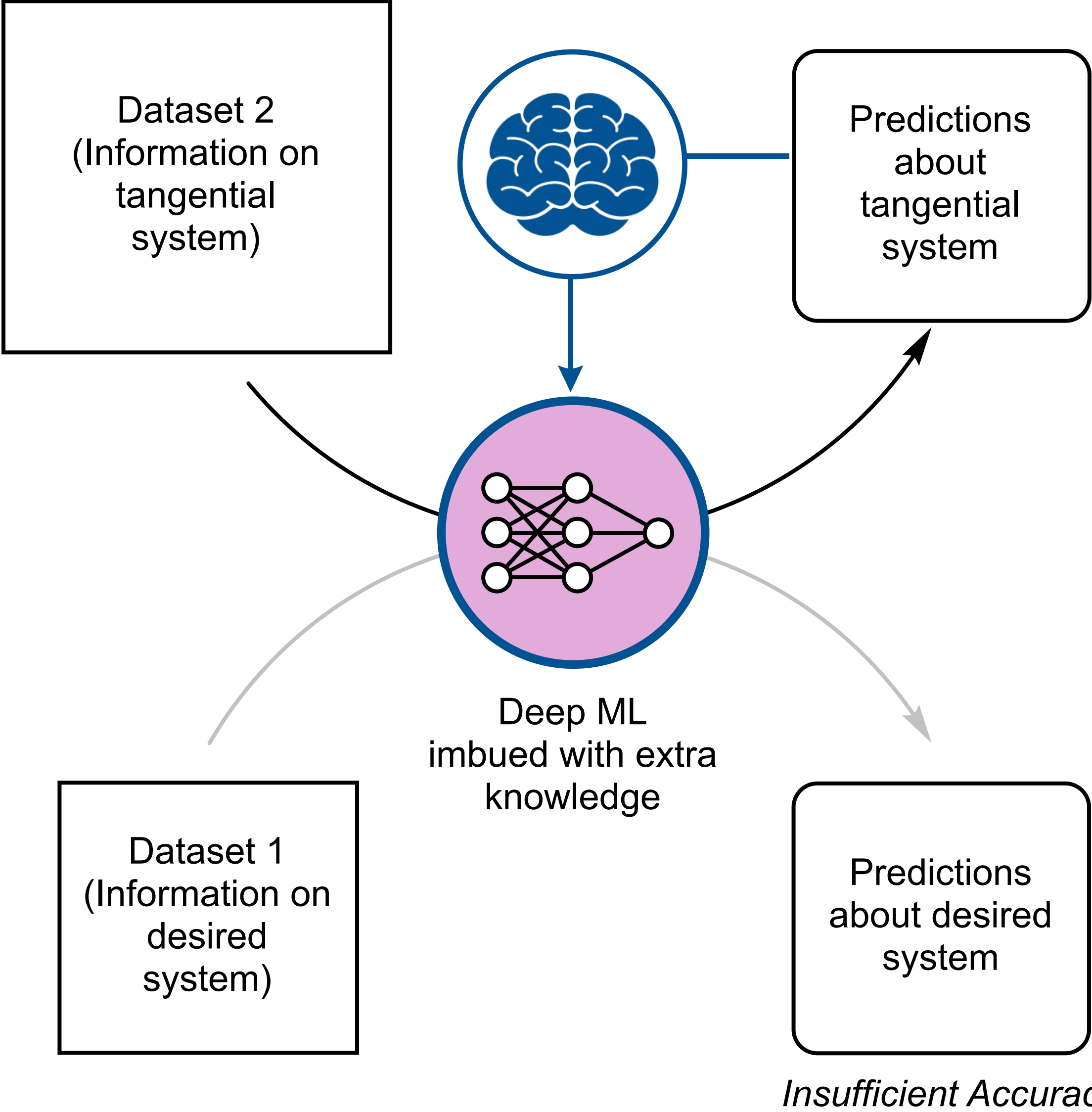
What Is Transfer Learning?



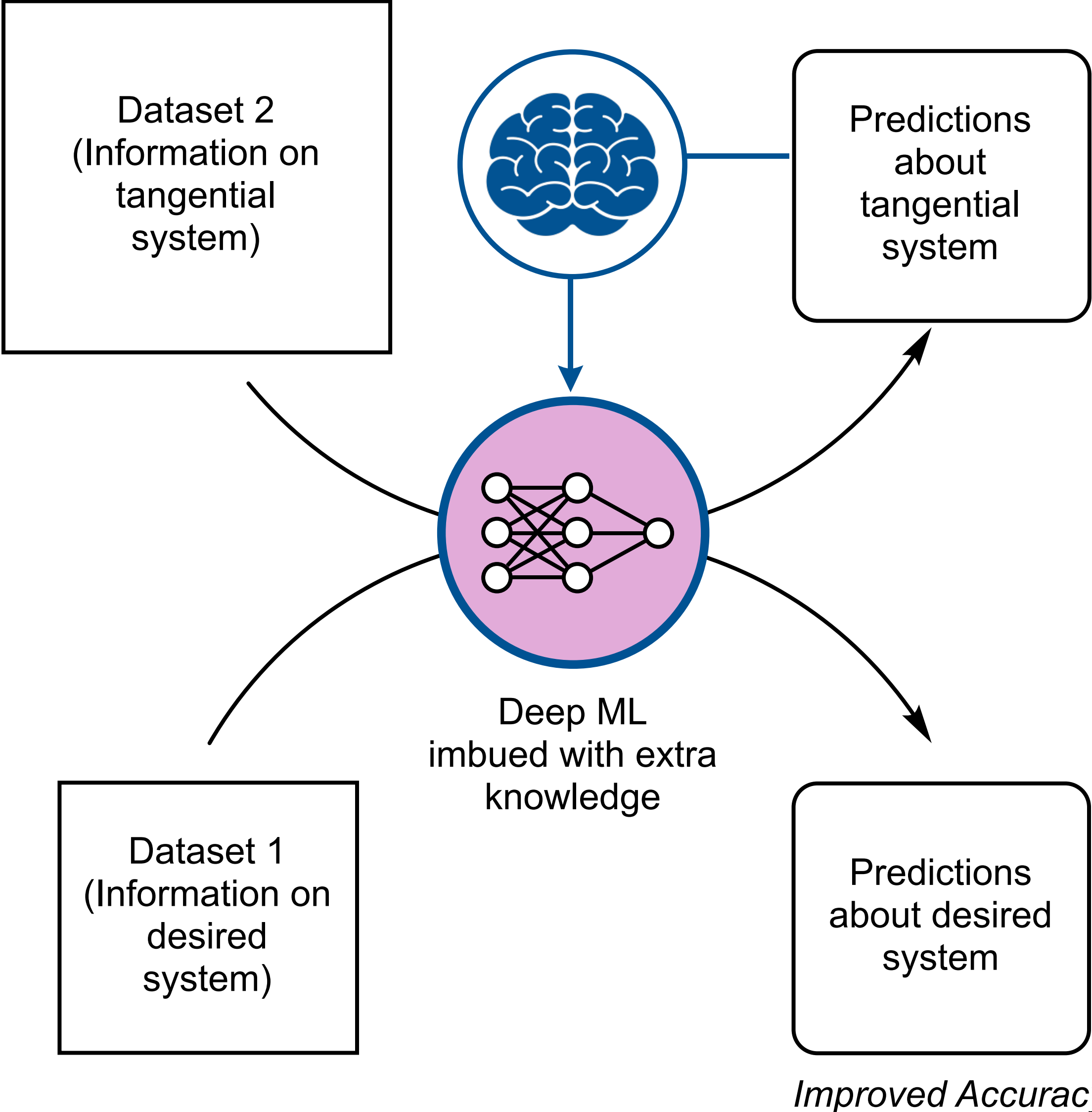
Predictions about desired system

Insufficient Accuracy

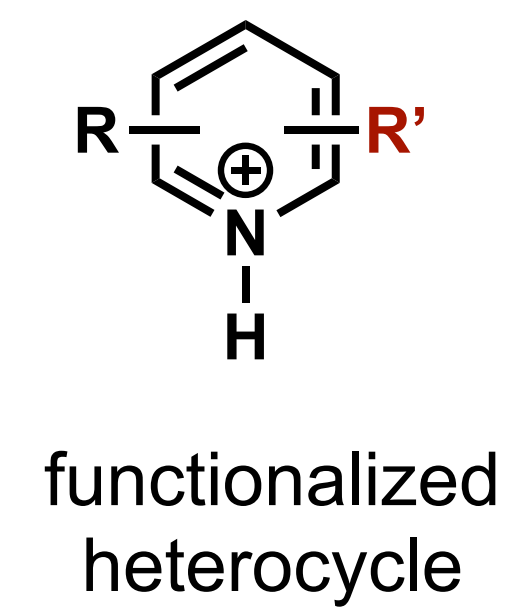
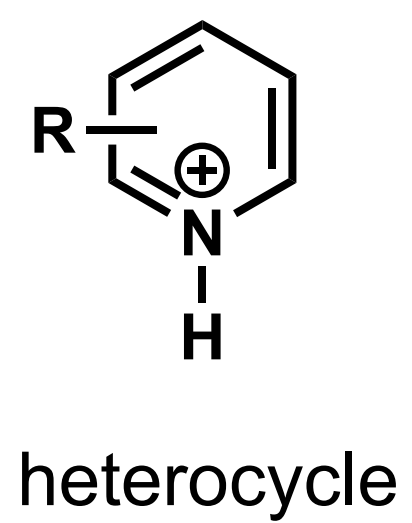
What Is Transfer Learning?



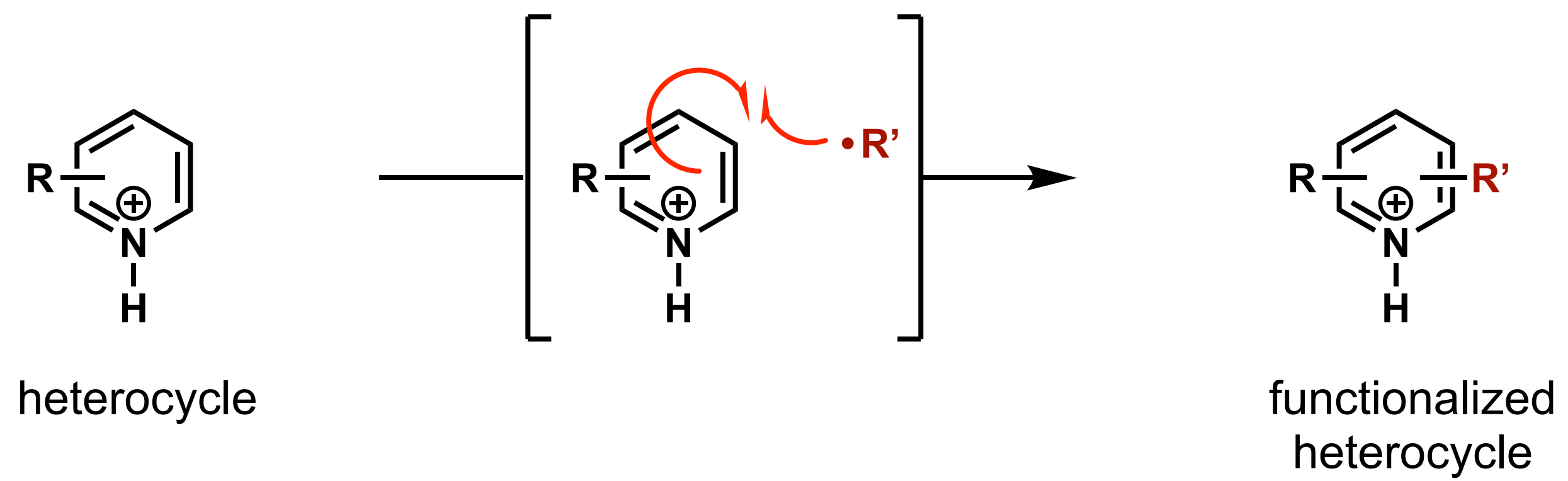
What Is Transfer Learning?



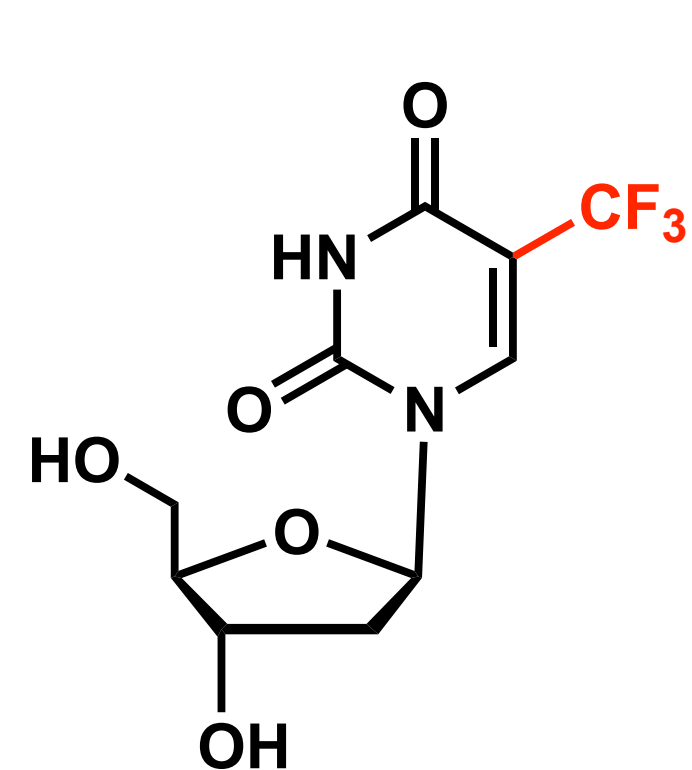
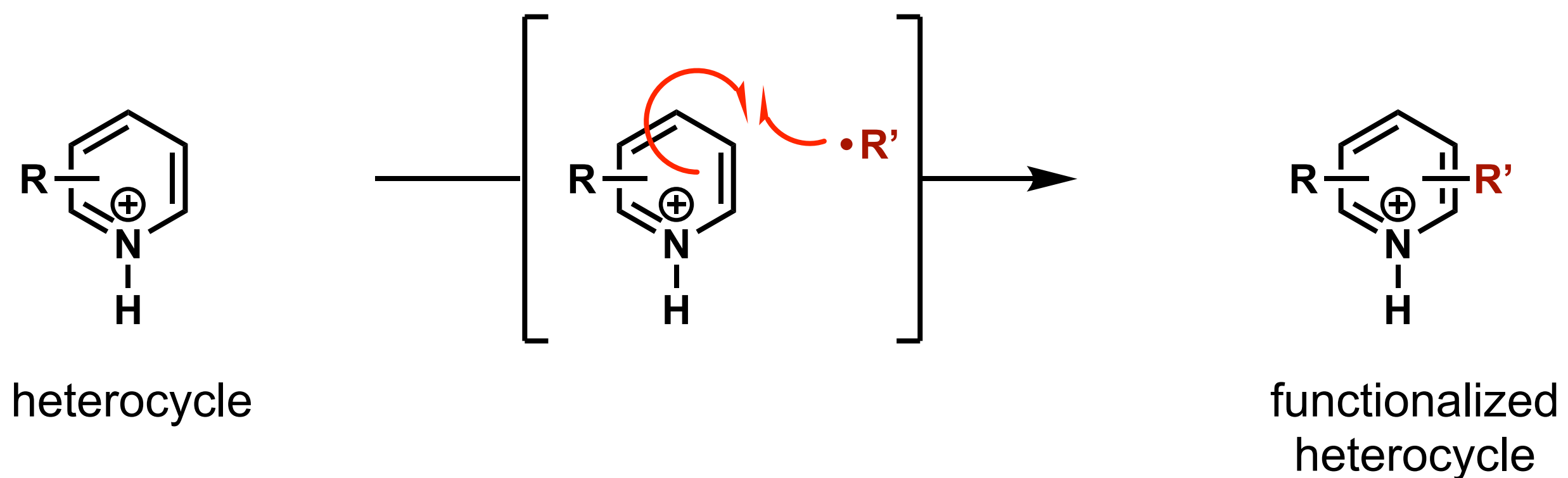
The Minisci Reaction



The Minisci Reaction

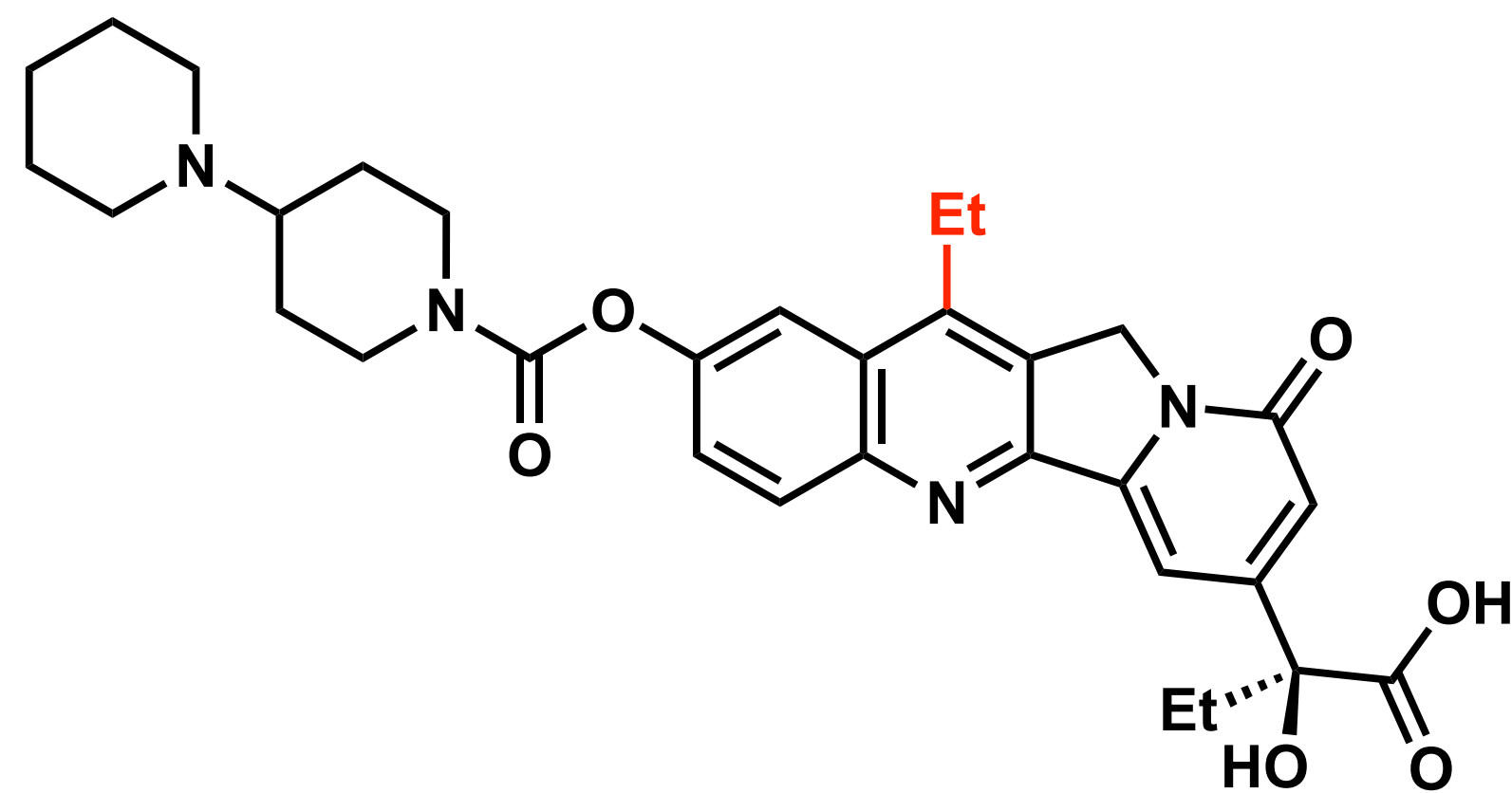


The Minisci Reaction



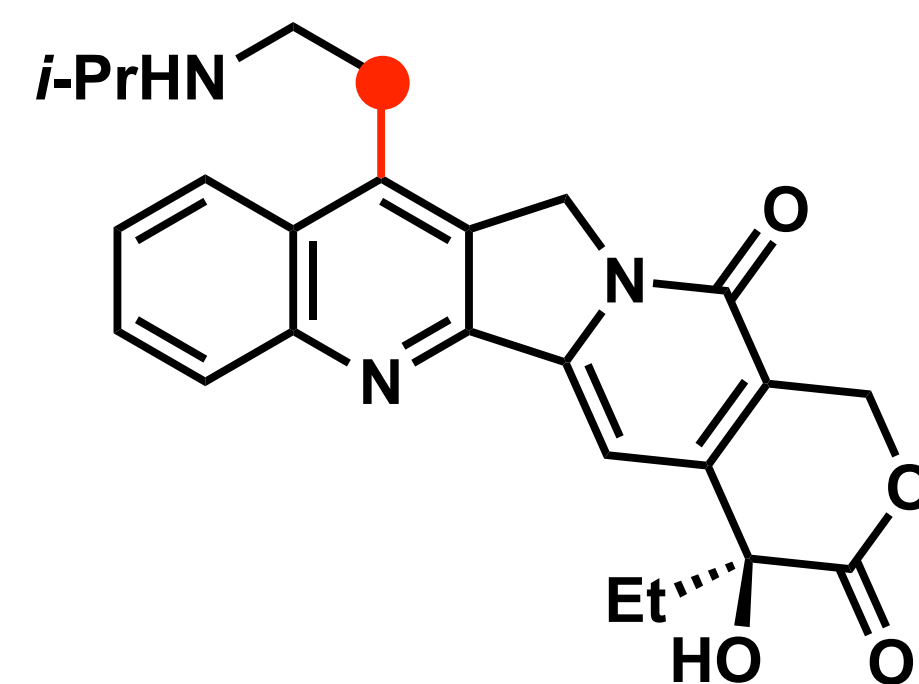
Viroptic

anti-viral eye drops



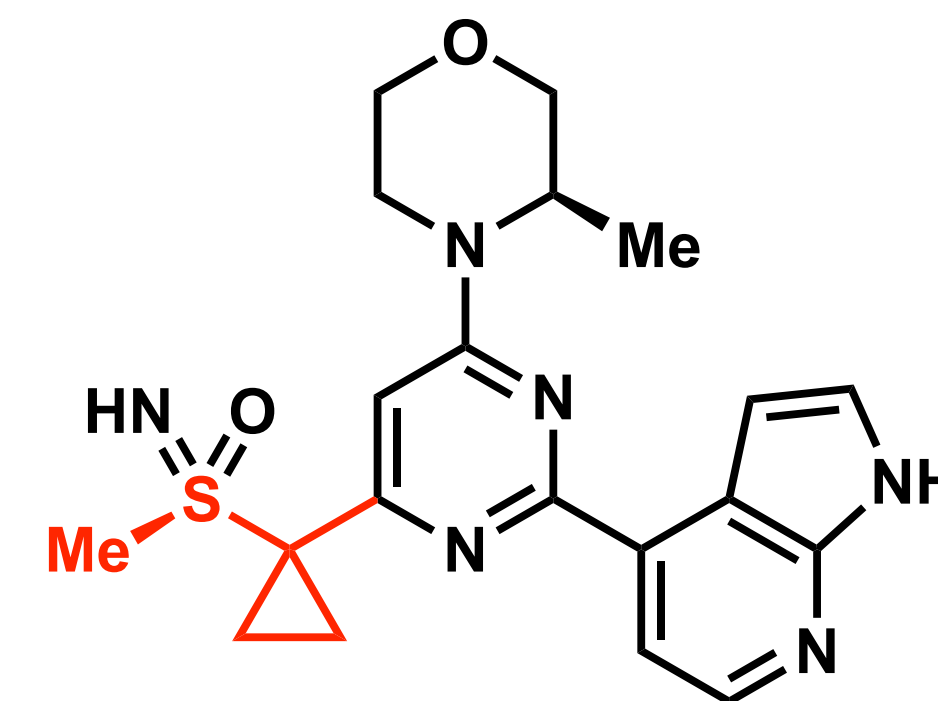
Irinotecan

colon, small-cell lung cancer



Belotecan

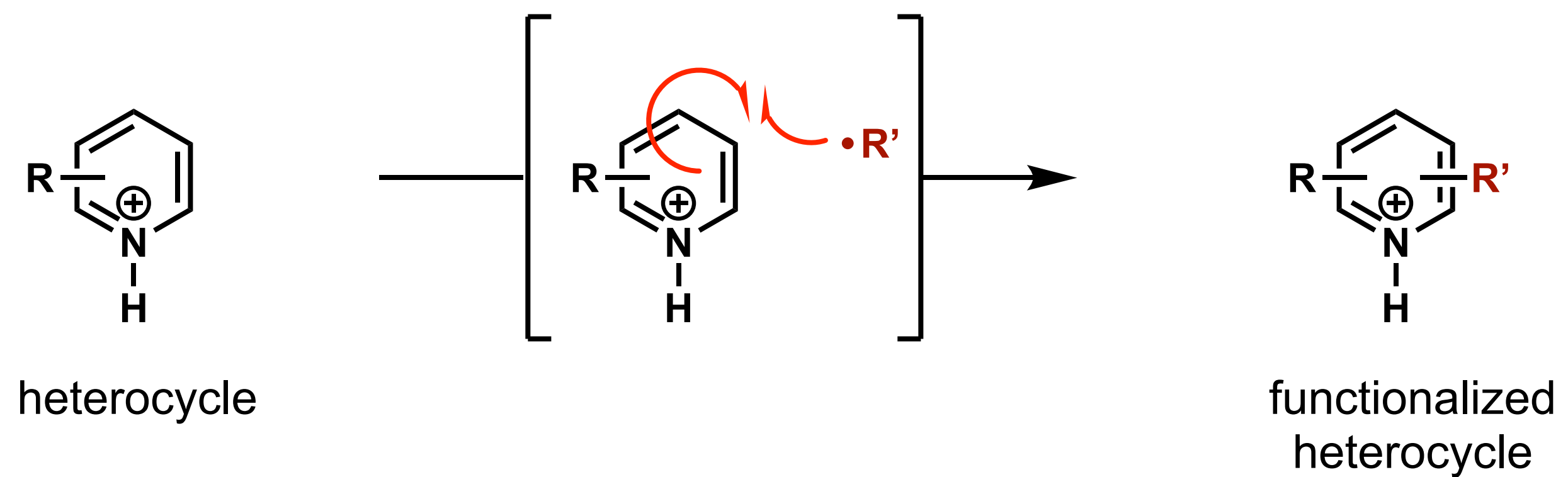
ovarian, small-cell lung cancer



Ceralasertib

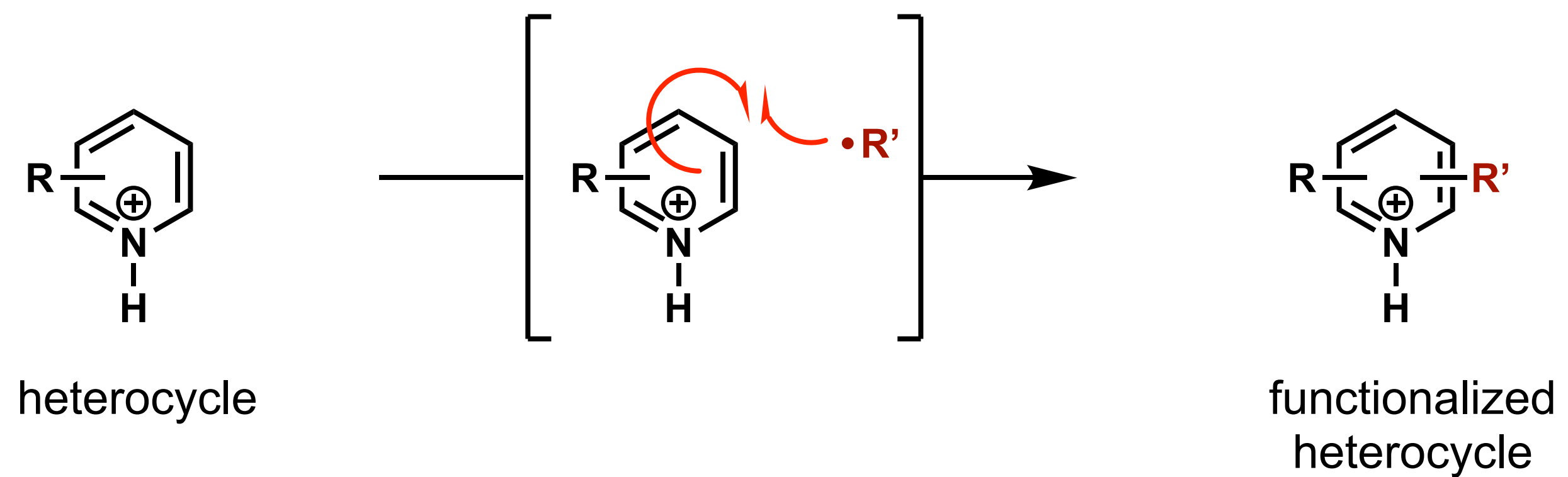
ovarian, small-cell lung, cervix cancer

Regioselectivity Factors of the Minisci Reaction



Electronics, sterics, and longevity of **•R'**

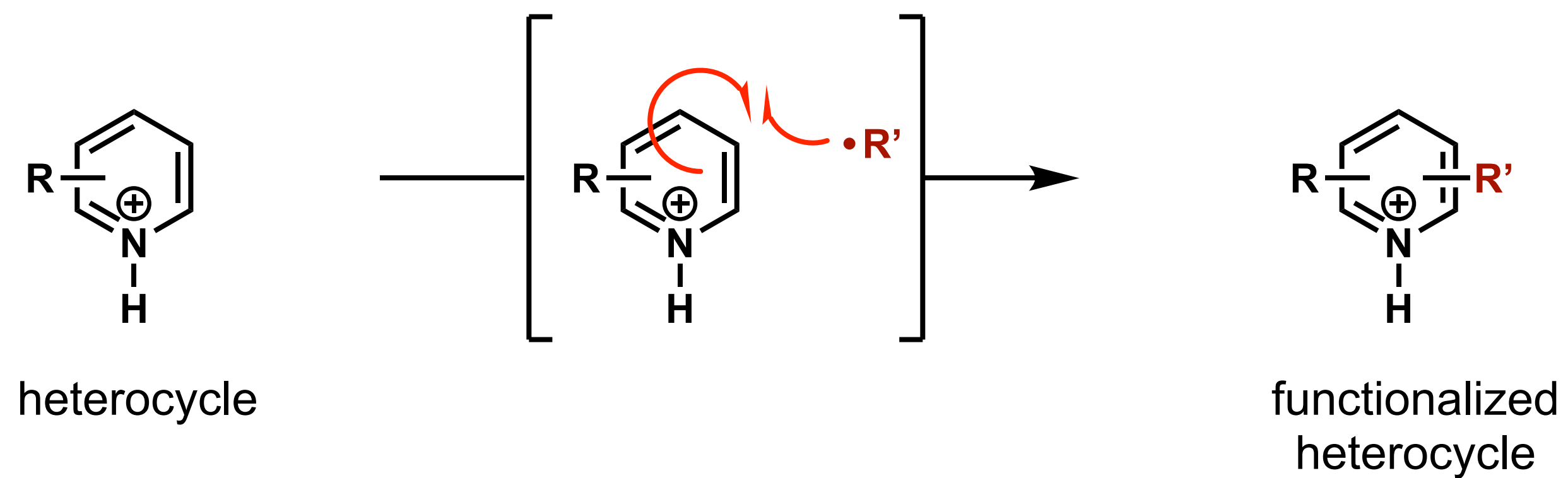
Regioselectivity Factors of the Minisci Reaction



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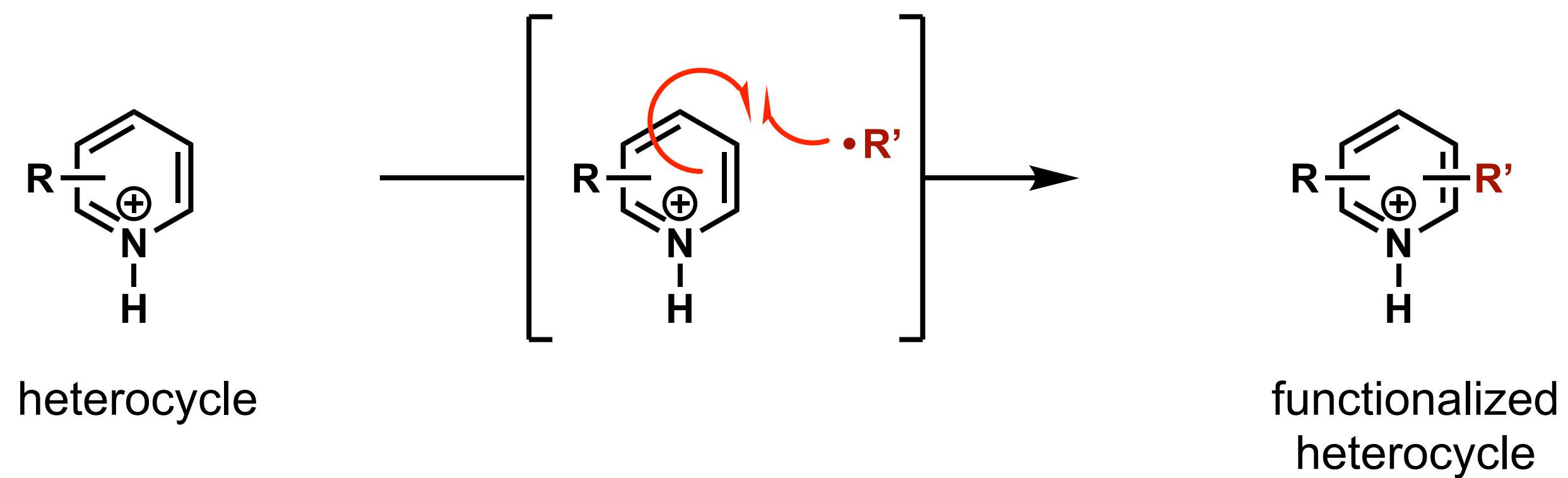
Electronics of heterocycle

Prediction of Regiochemical Outcomes of the Minisci Reaction



DFT-derived Fukui reactivity indices are ~90% accurate.

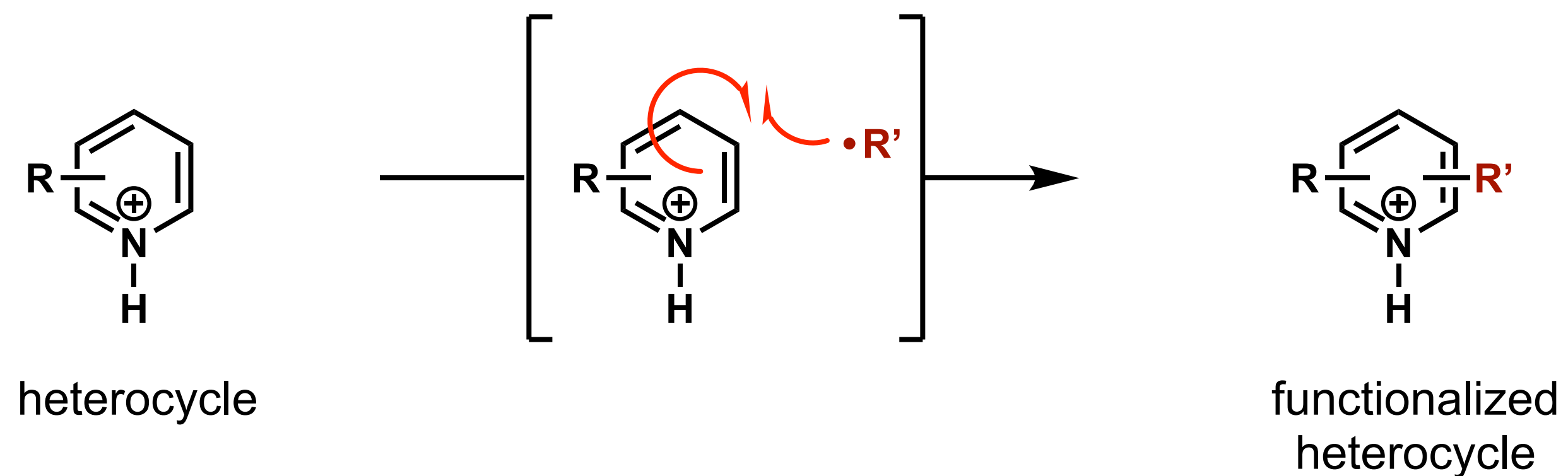
Prediction of Regiochemical Outcomes of the Minisci Reaction



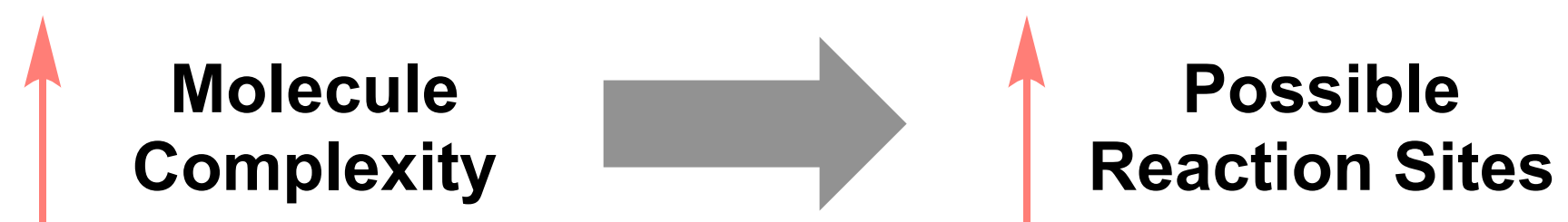
DFT-derived Fukui reactivity indices are ~90% accurate.

↑
**Molecule
Complexity**

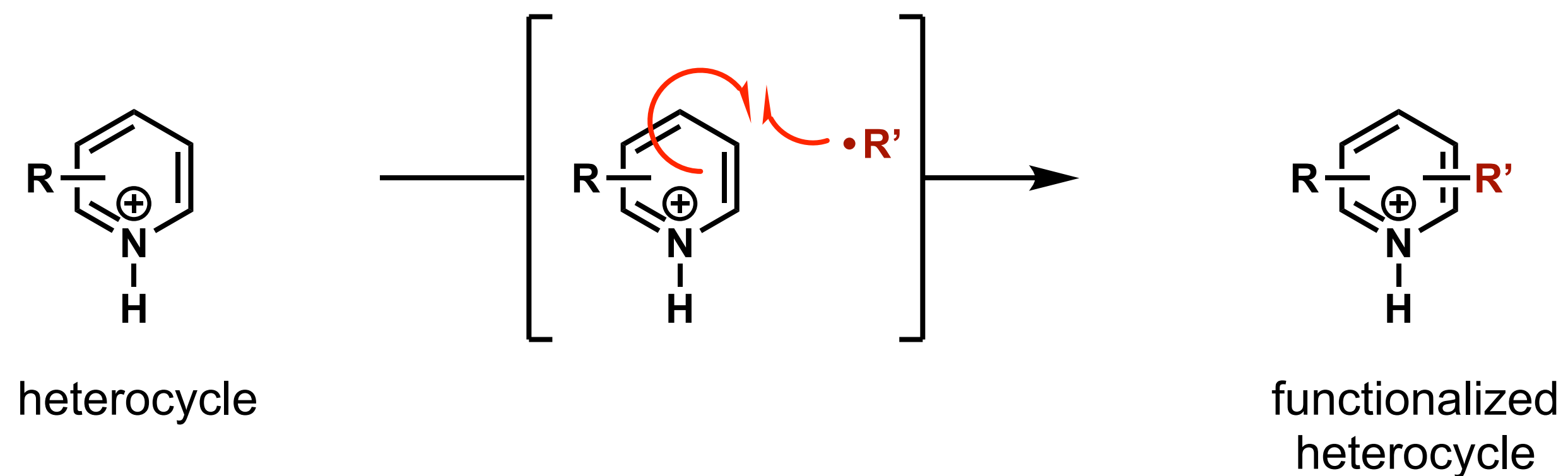
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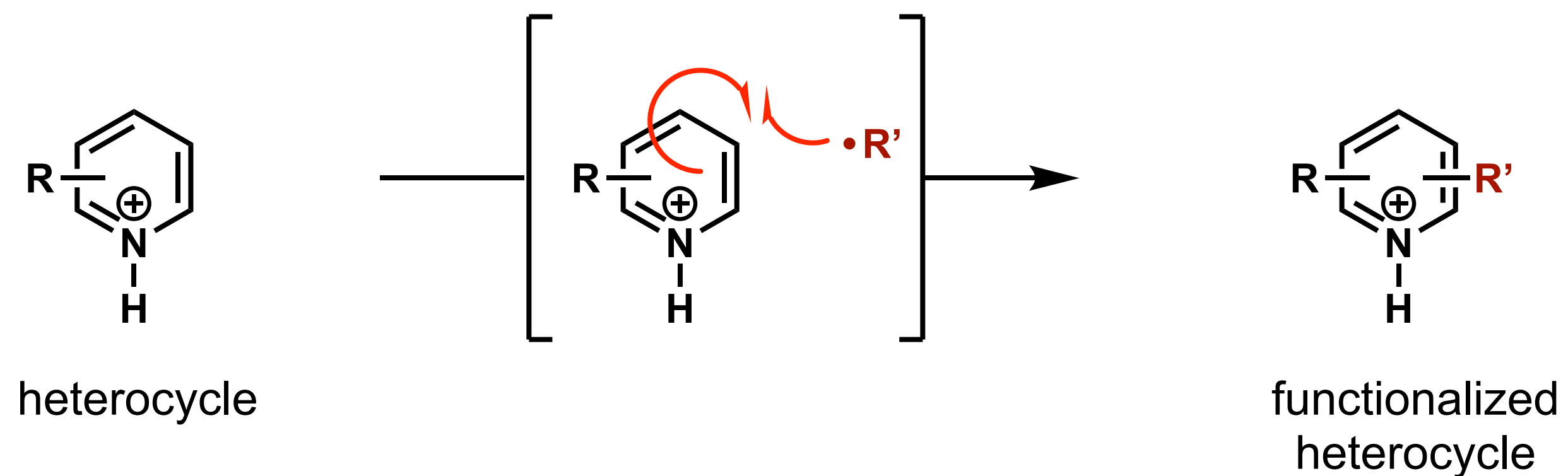
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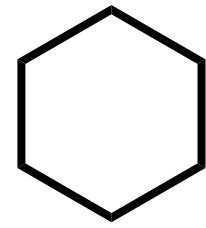


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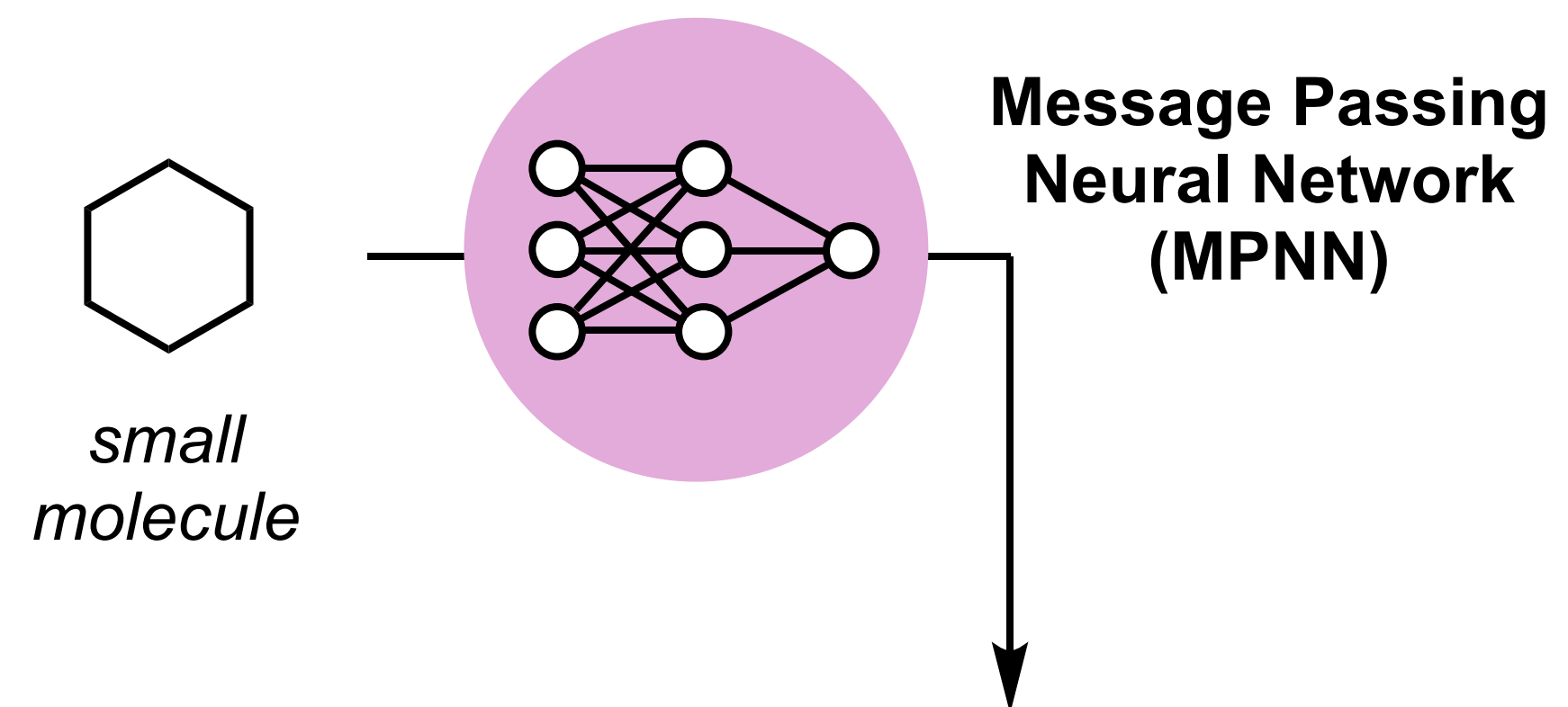
Can machine learning provide some improvement?

The Big Idea

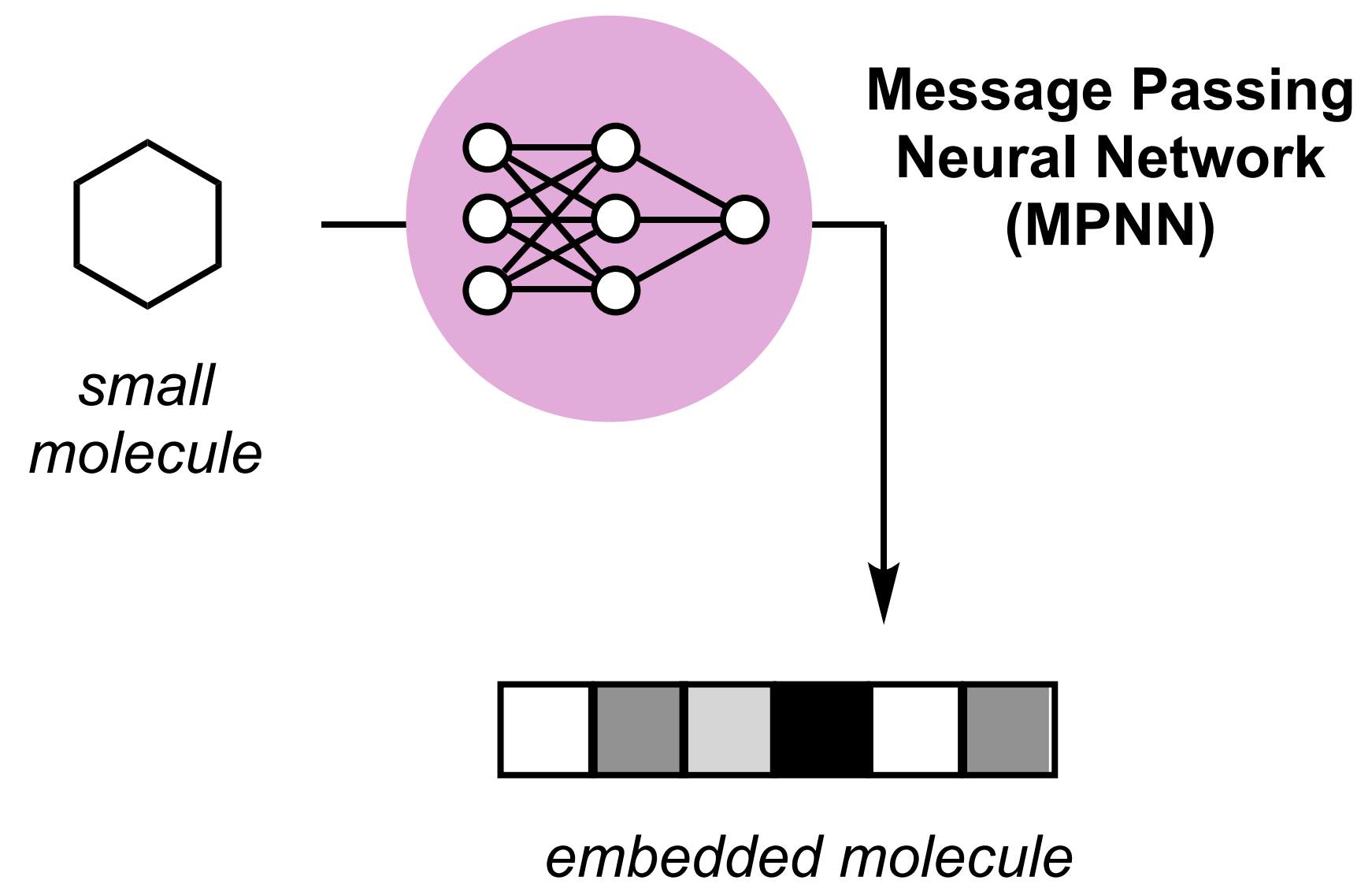


*small
molecule*

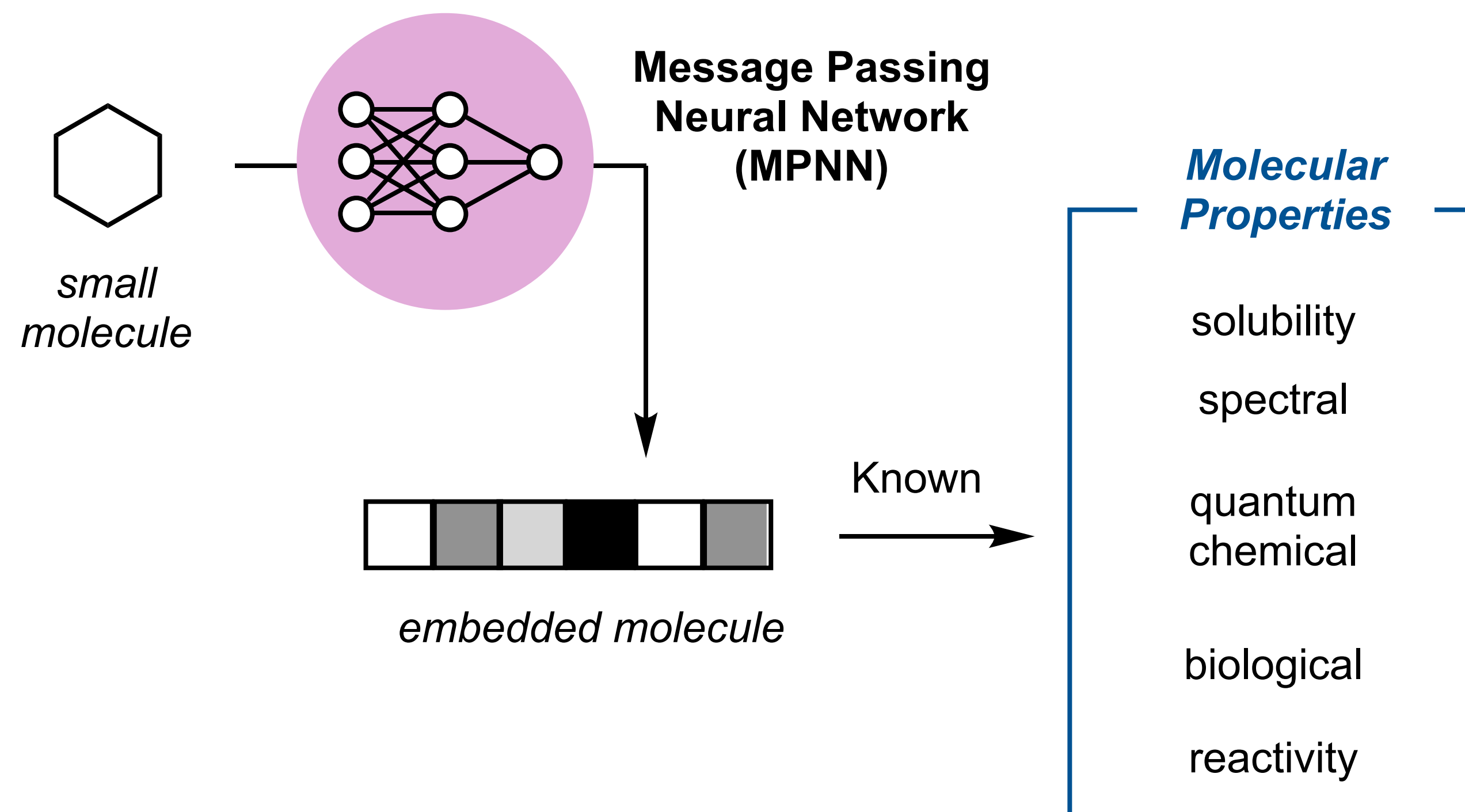
The Big Idea

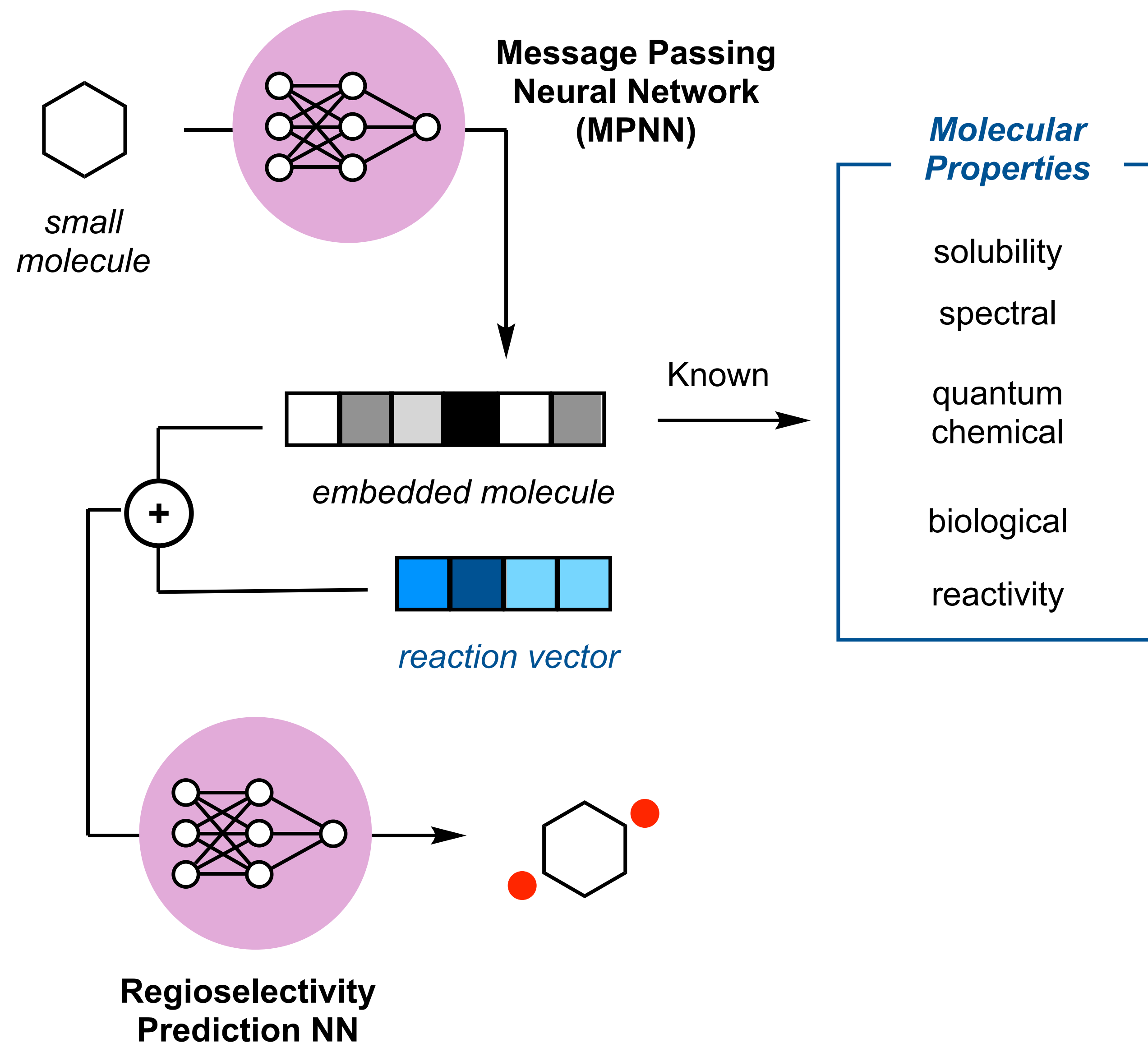


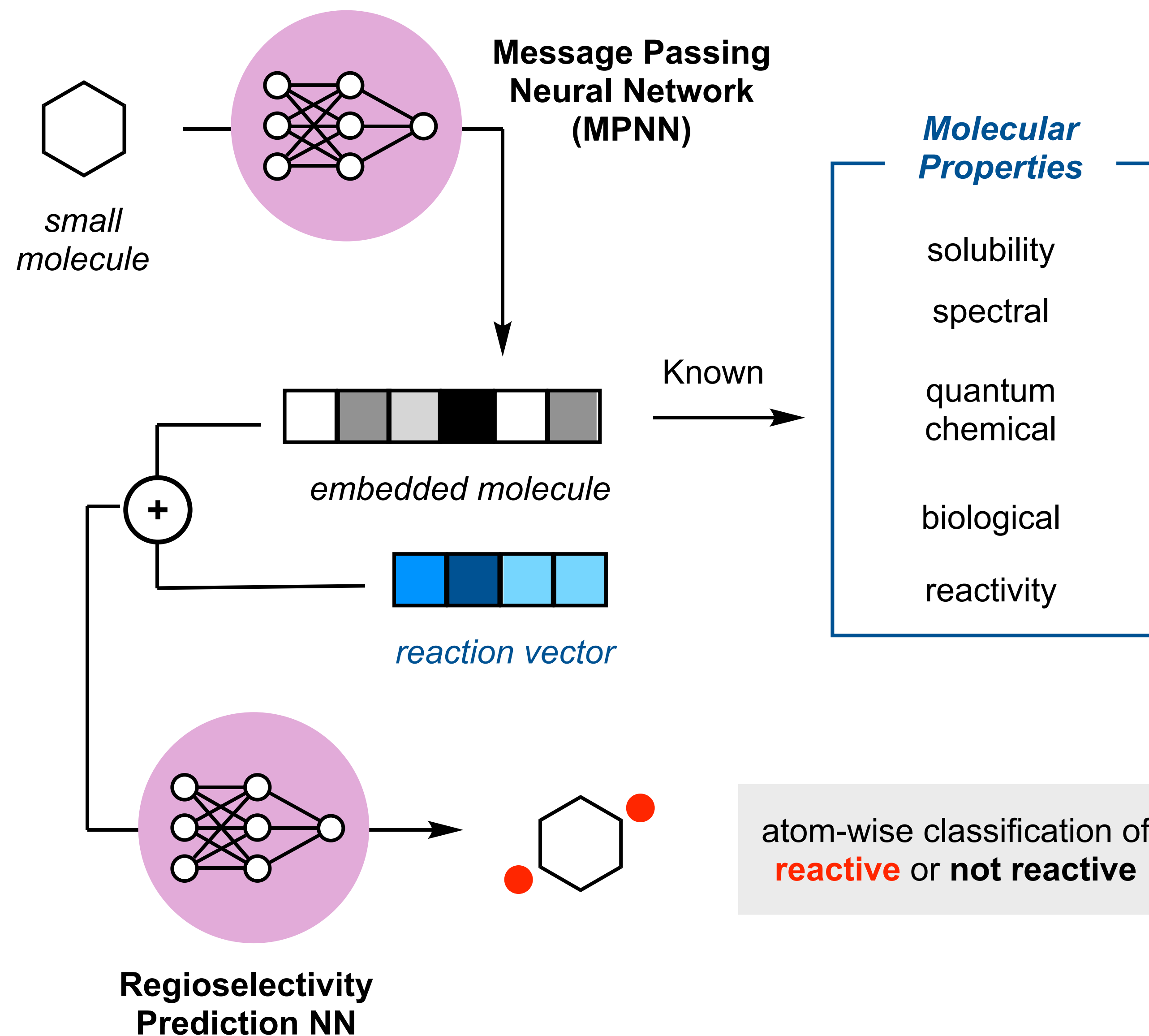
The Big Idea



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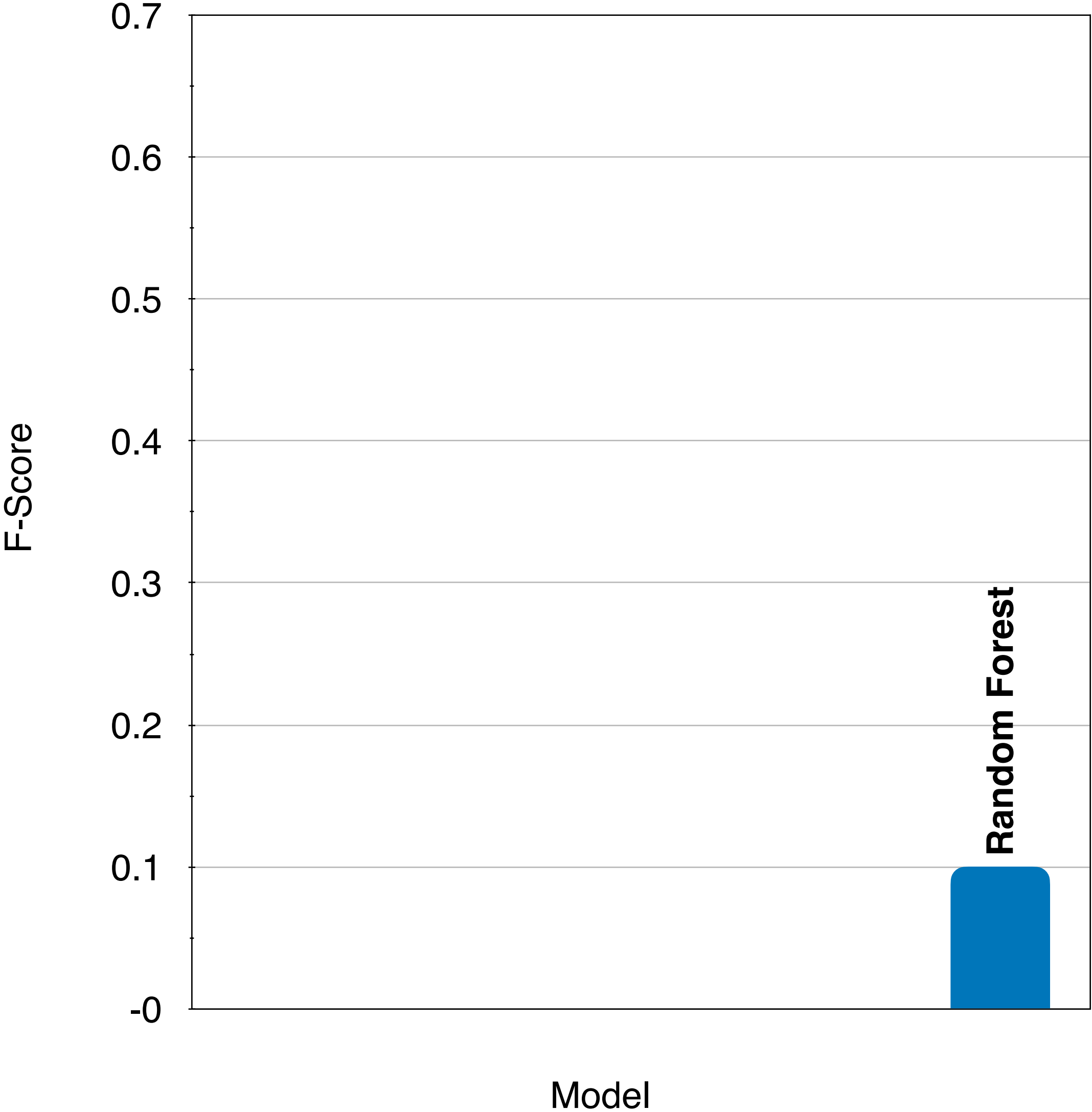




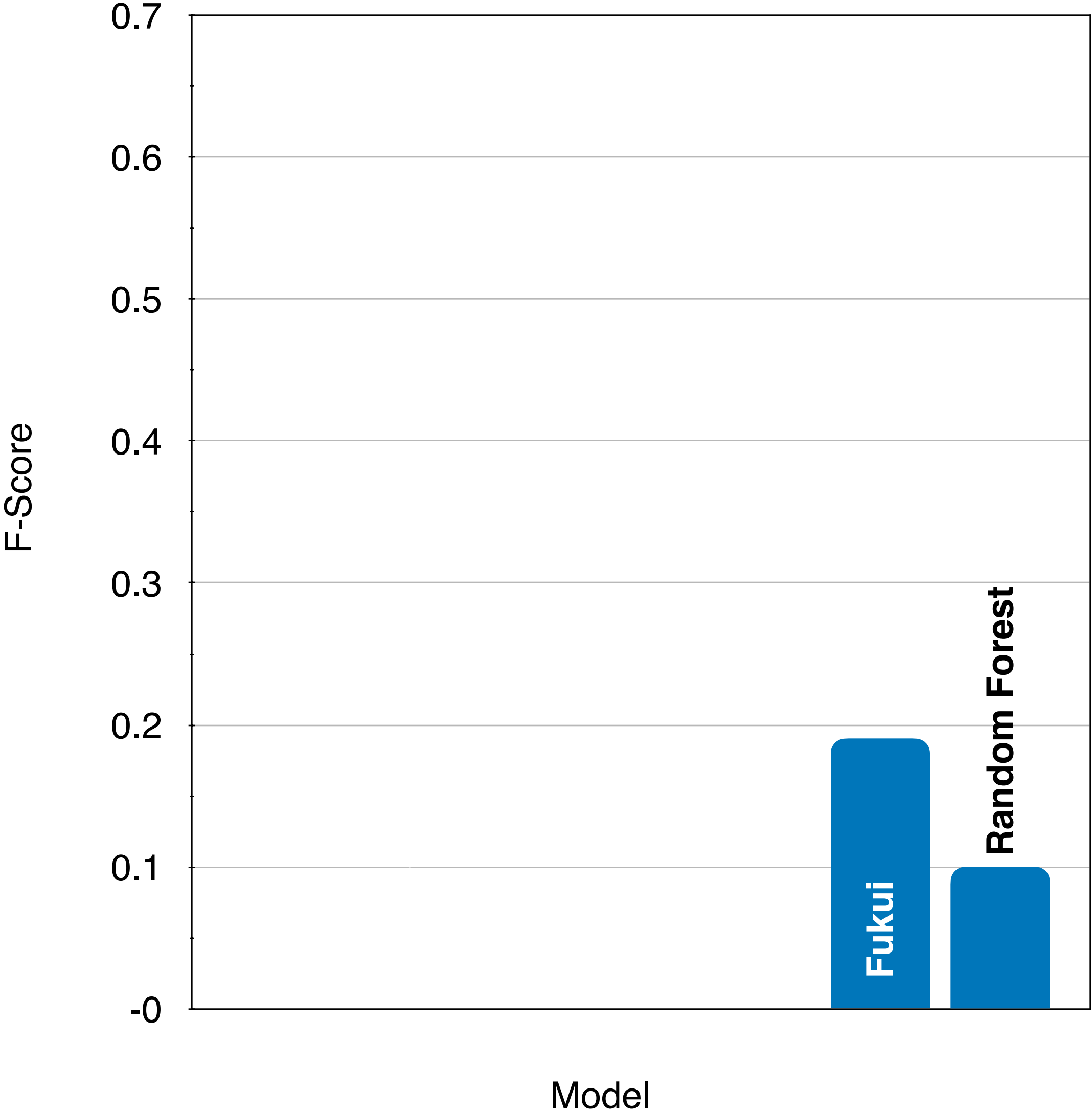


atom-wise classification of
reactive or **not reactive**

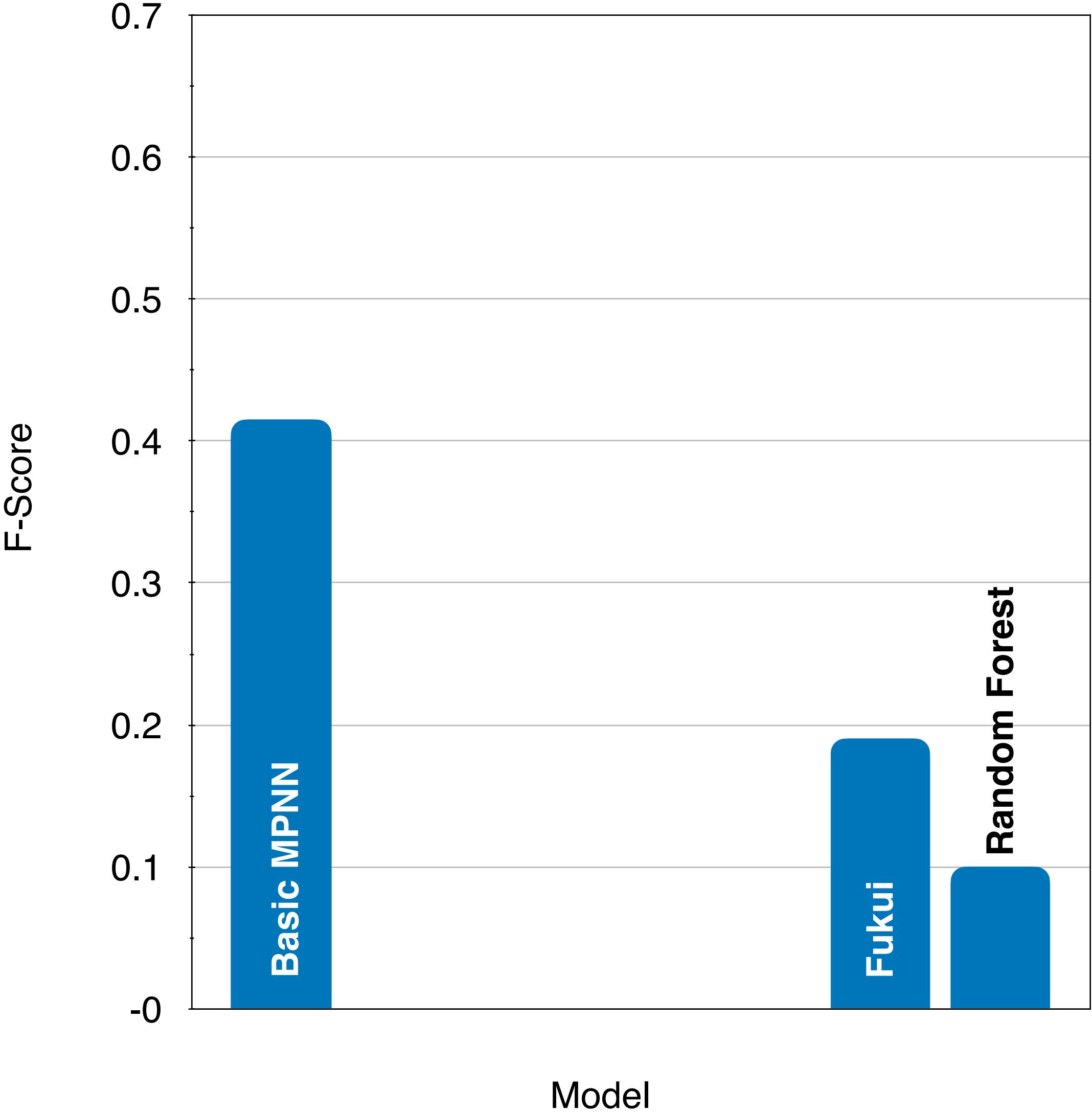
Model Accuracy (F-Score)



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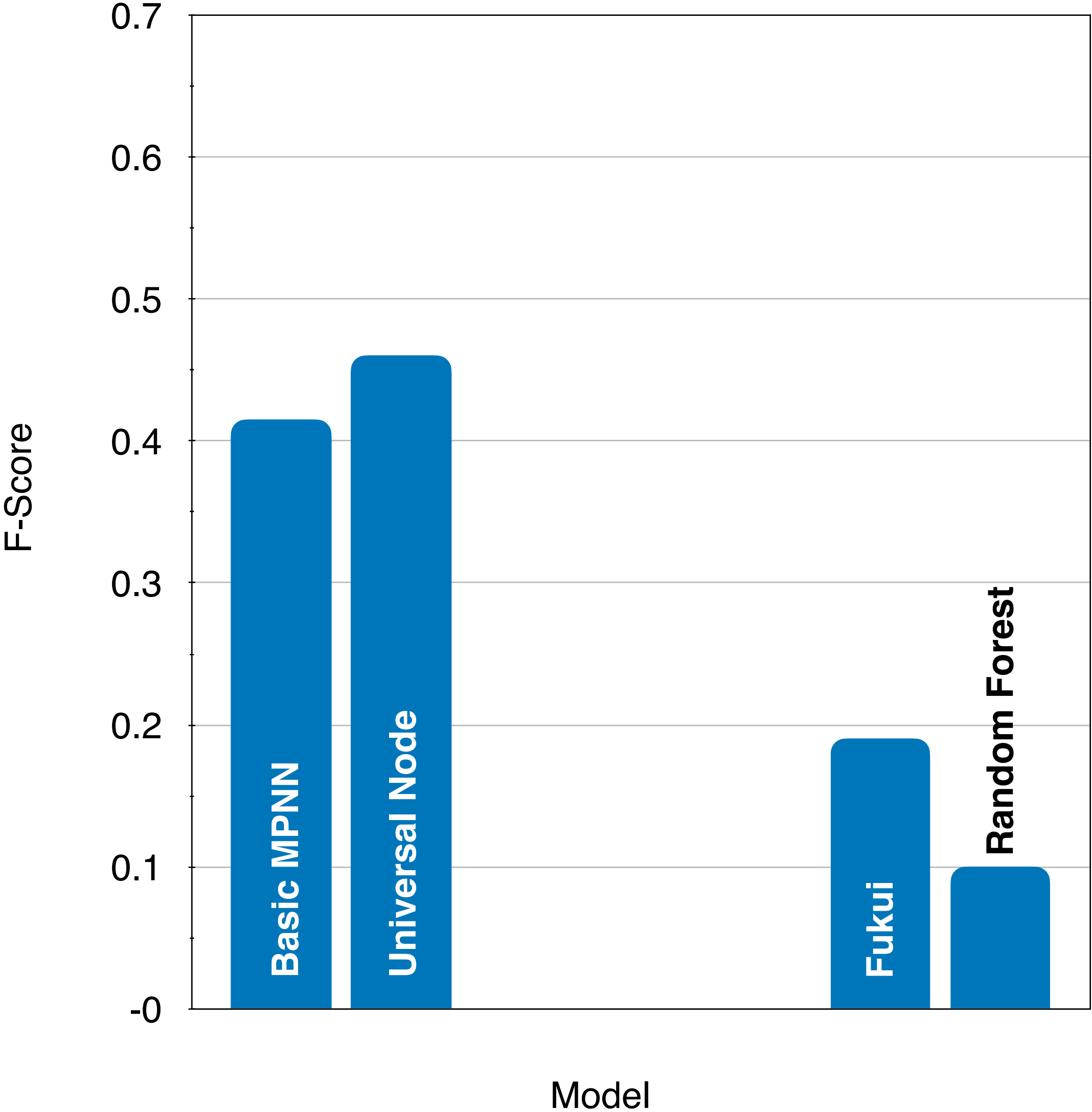


Model Accuracy (F-Score)

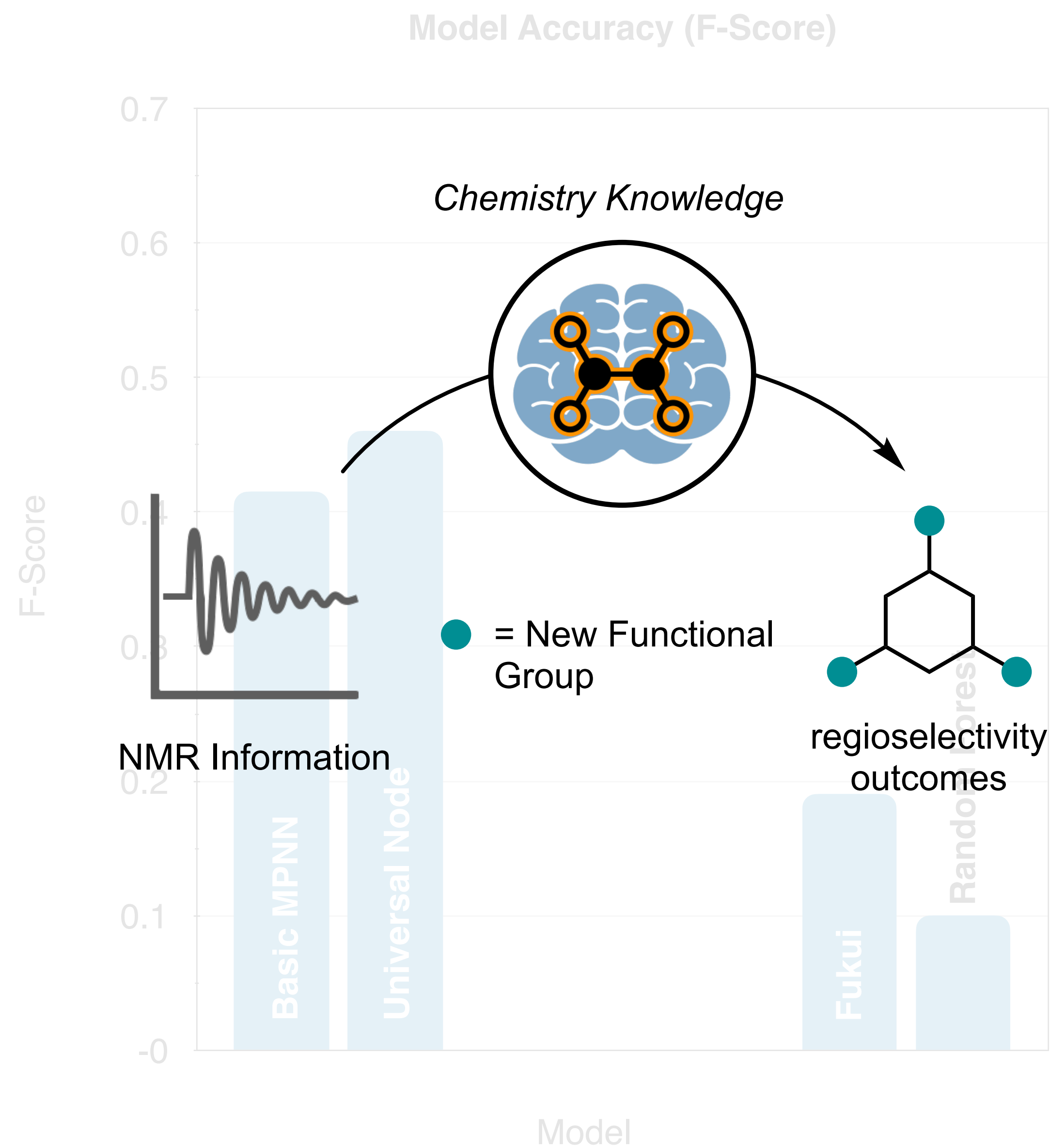


A Modest Improvement

Model Accuracy (F-Score)

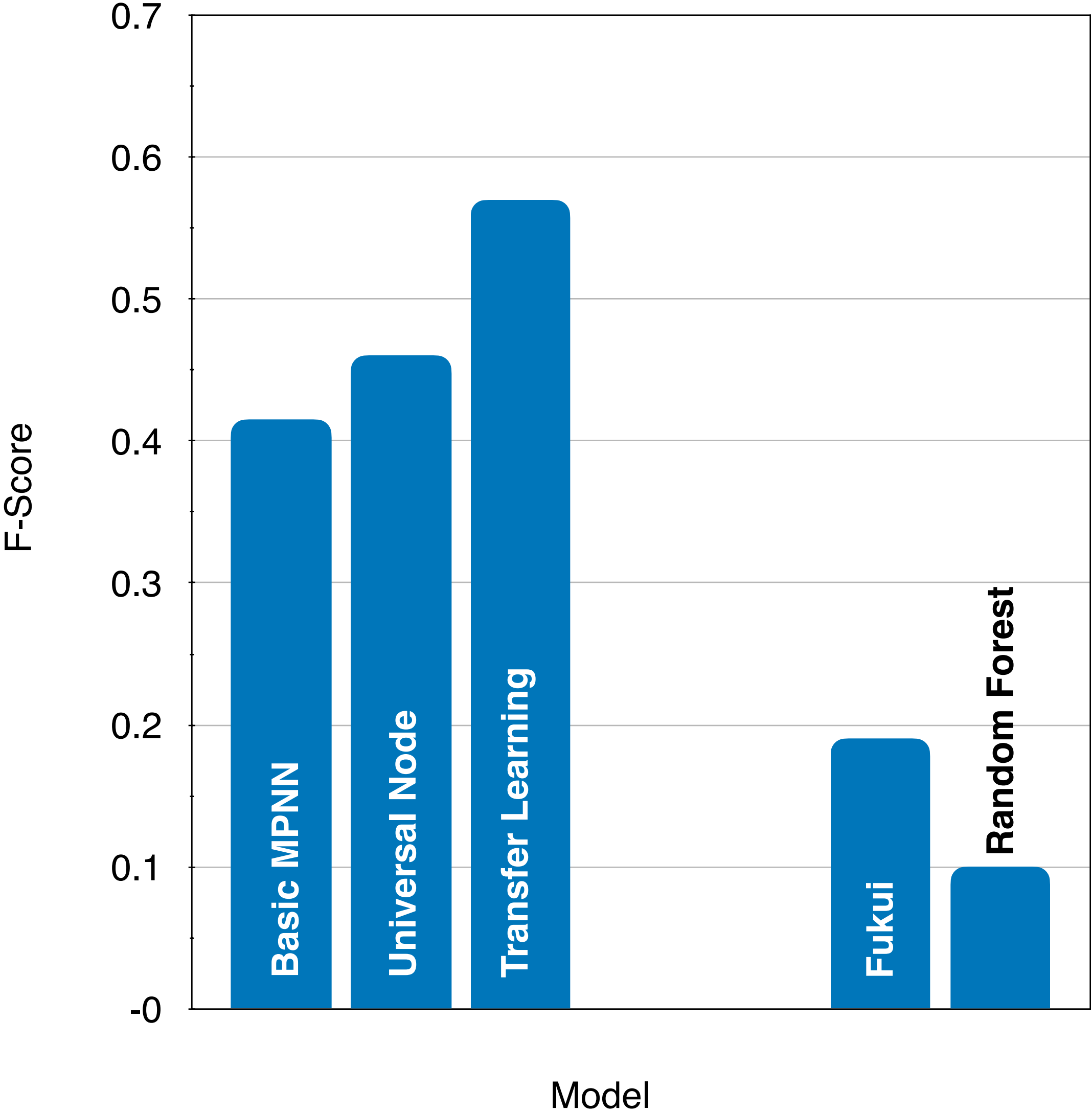


A Modest Improvement



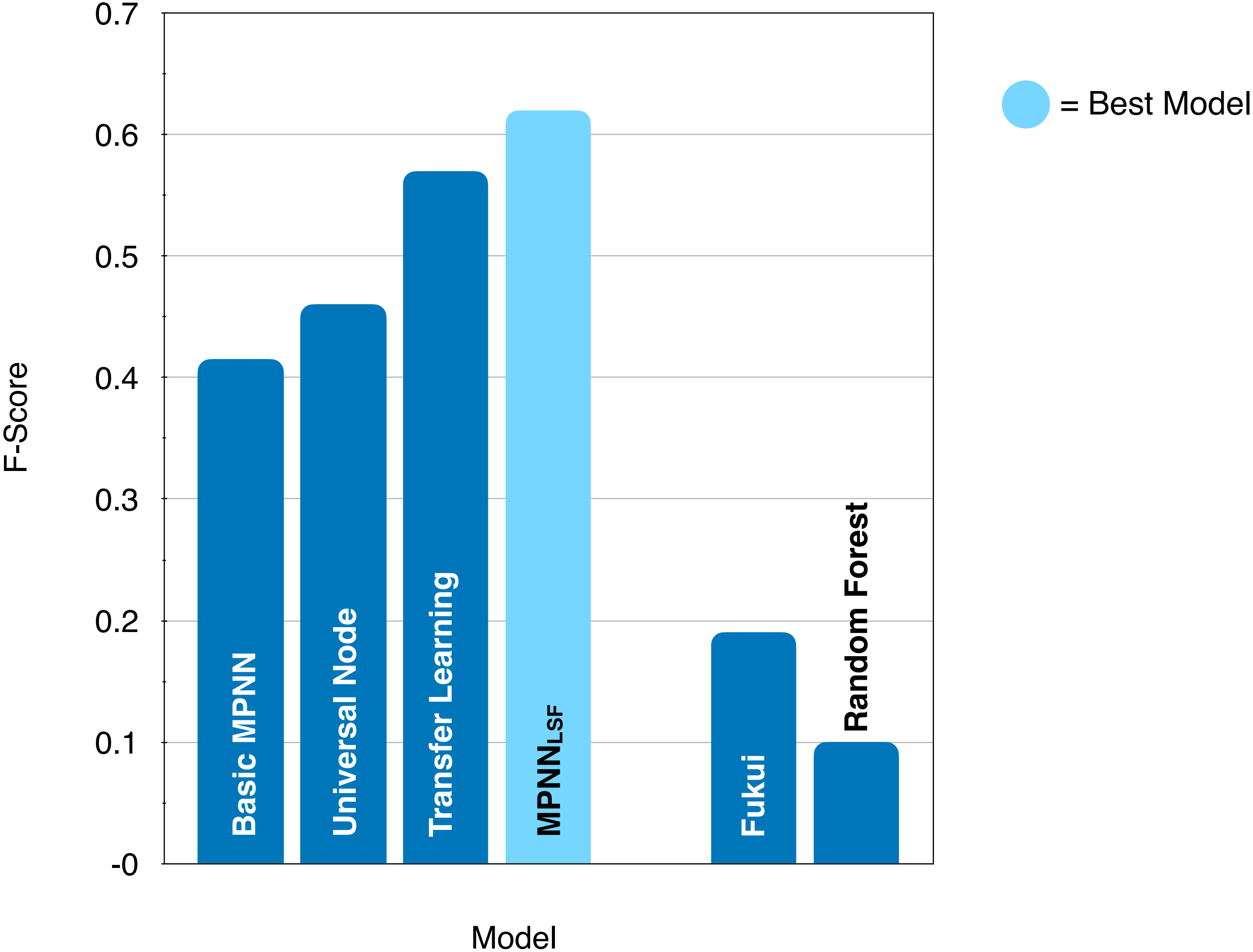
Significant Improvement!

Model Accuracy (F-Score)

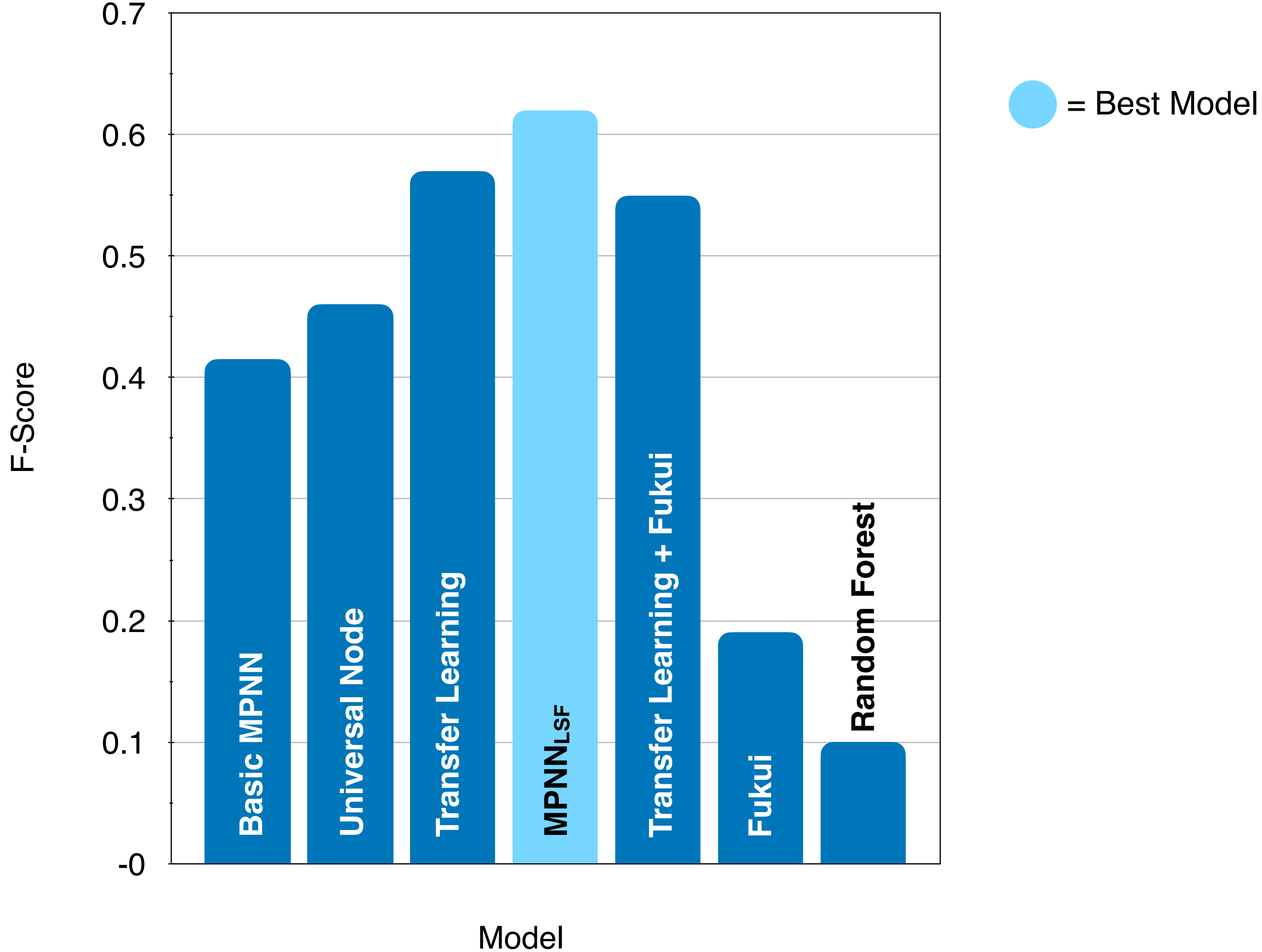


Significant Improvement!

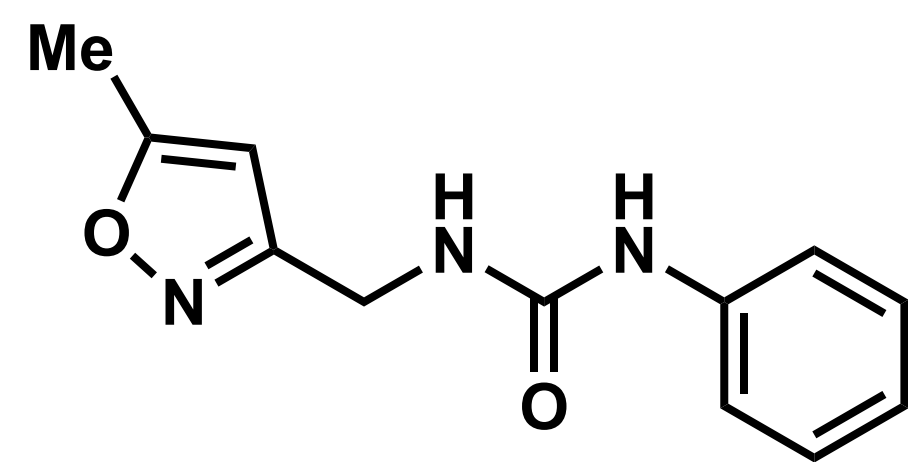
Model Accuracy (F-Score)



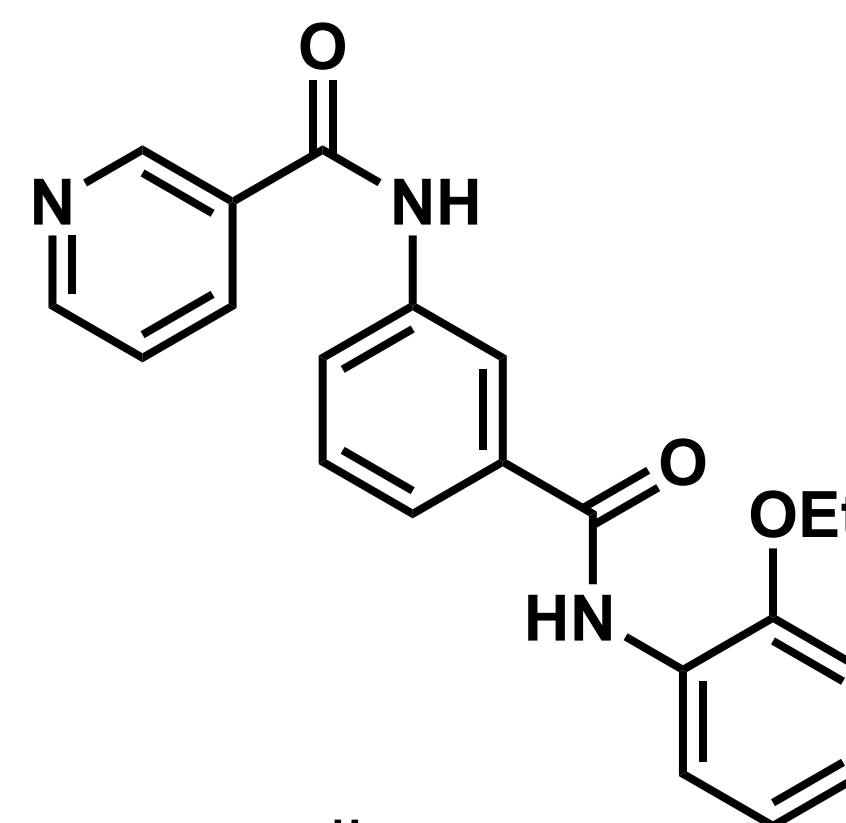
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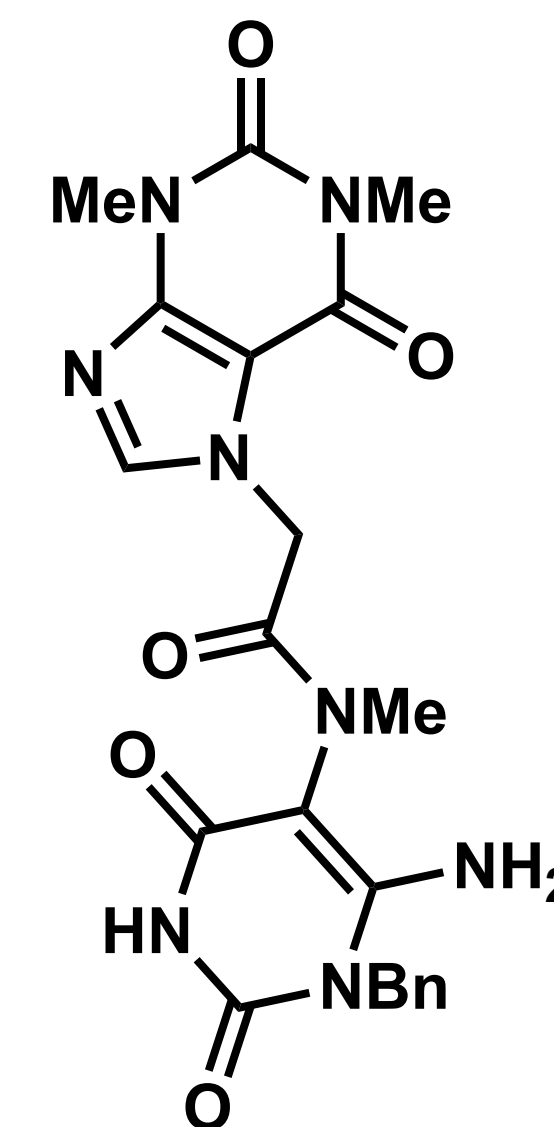
How Do We Perform on Completely Unseen Molecules?



small



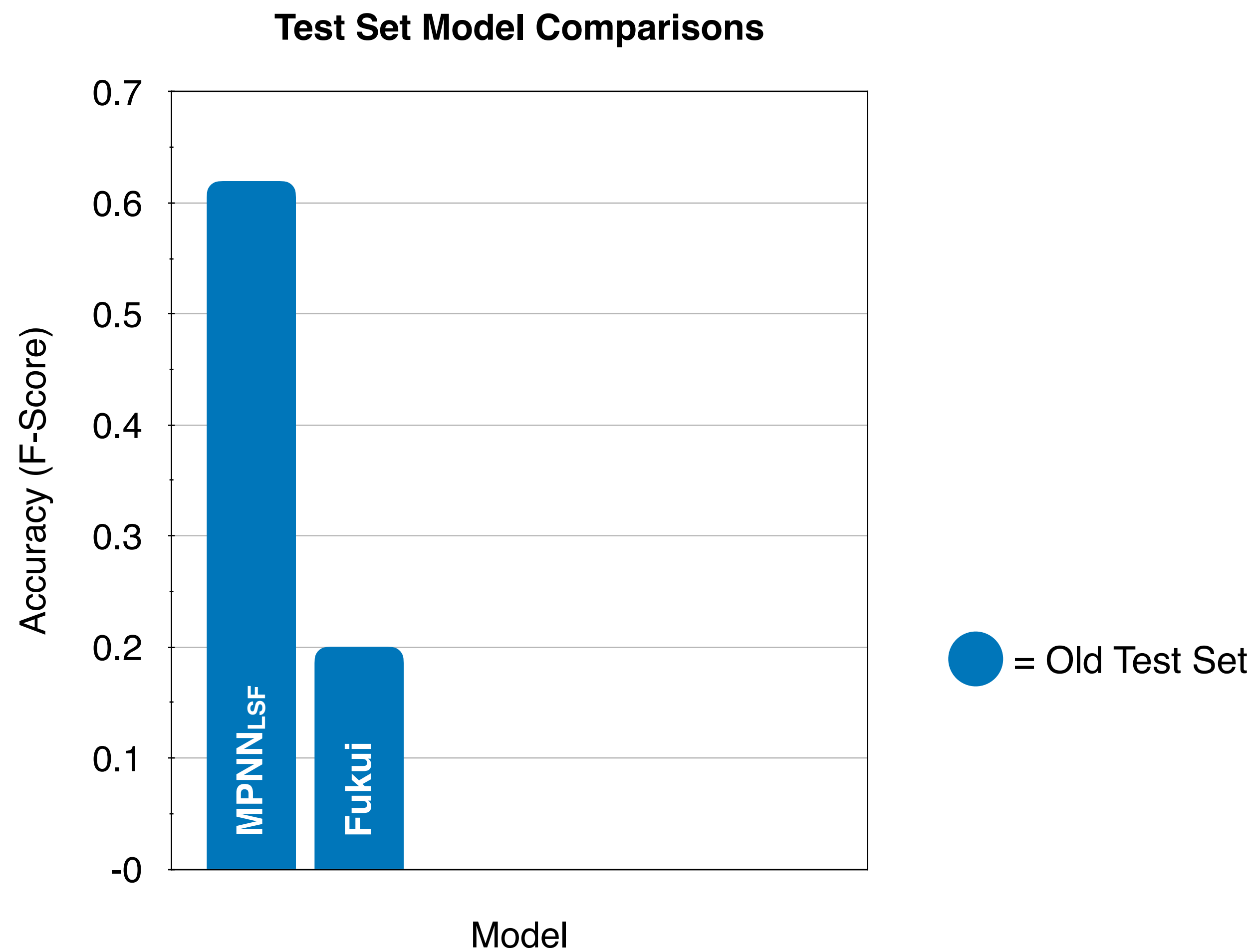
medium



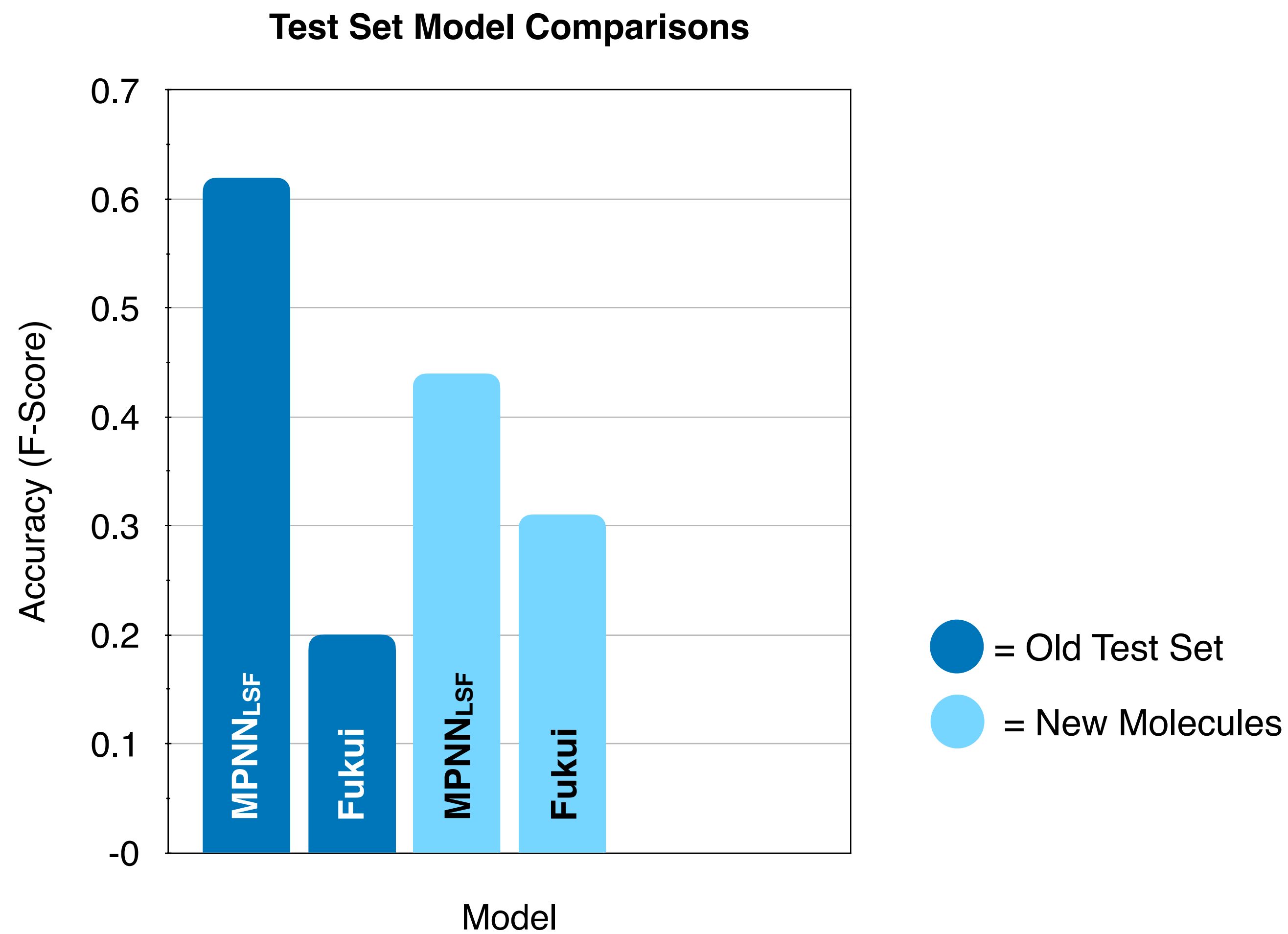
large

How do we do in a real-life scenario?

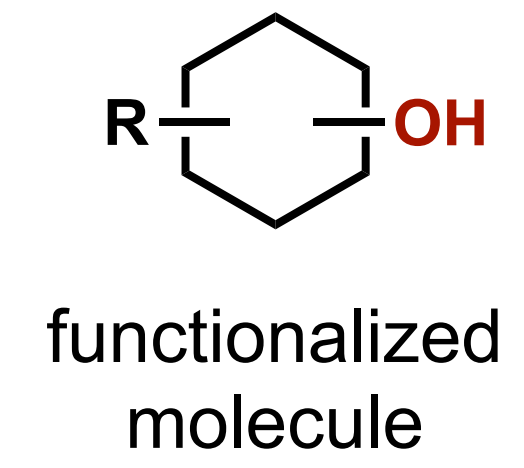
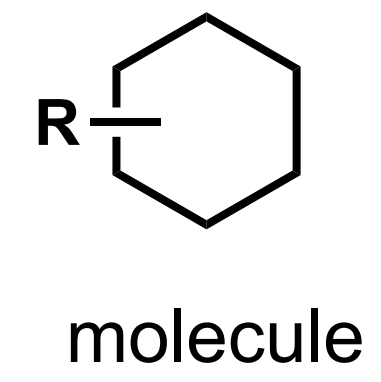
How Do We Perform on Completely Unseen Molecules?



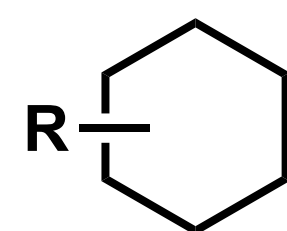
How Do We Perform on Completely Unseen Molecules?



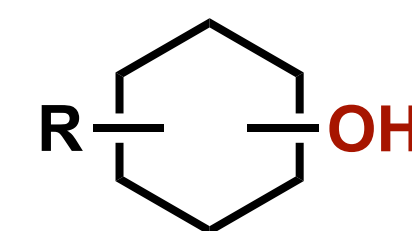
A Different Reaction Altogether



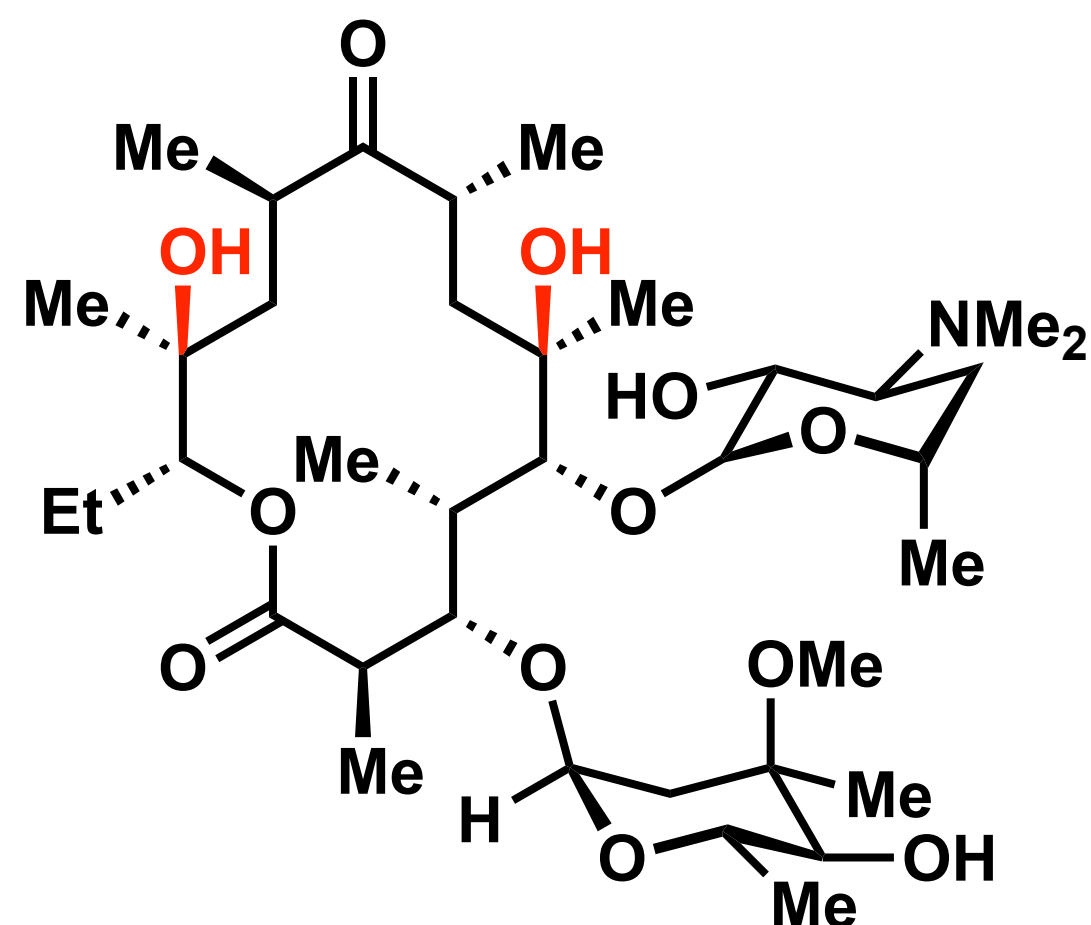
A Different Reaction Altogether



molecule

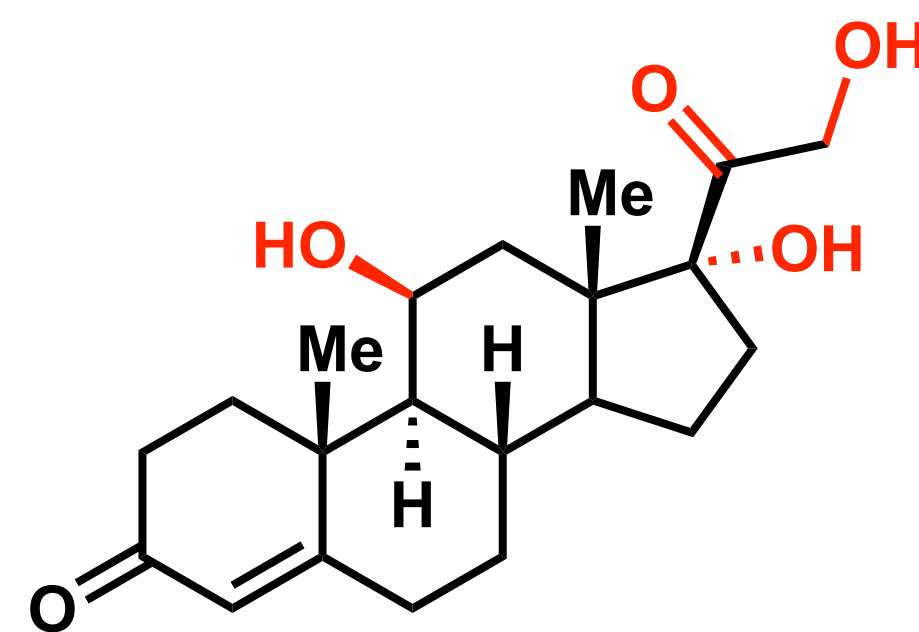


functionalized
molecule



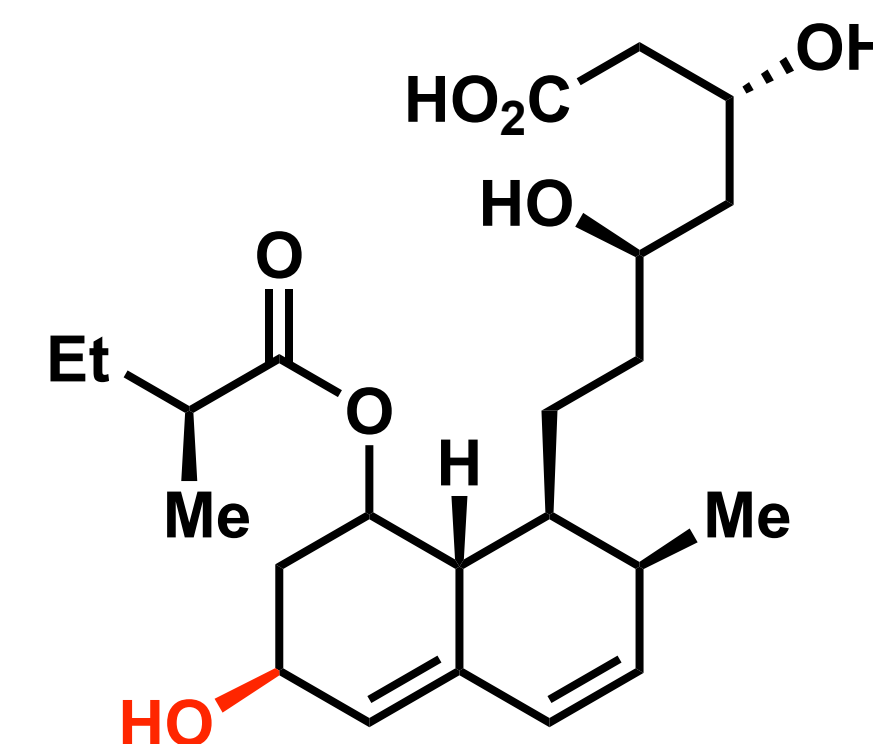
erythromycin

common anti-bacterial



hydrocortisone

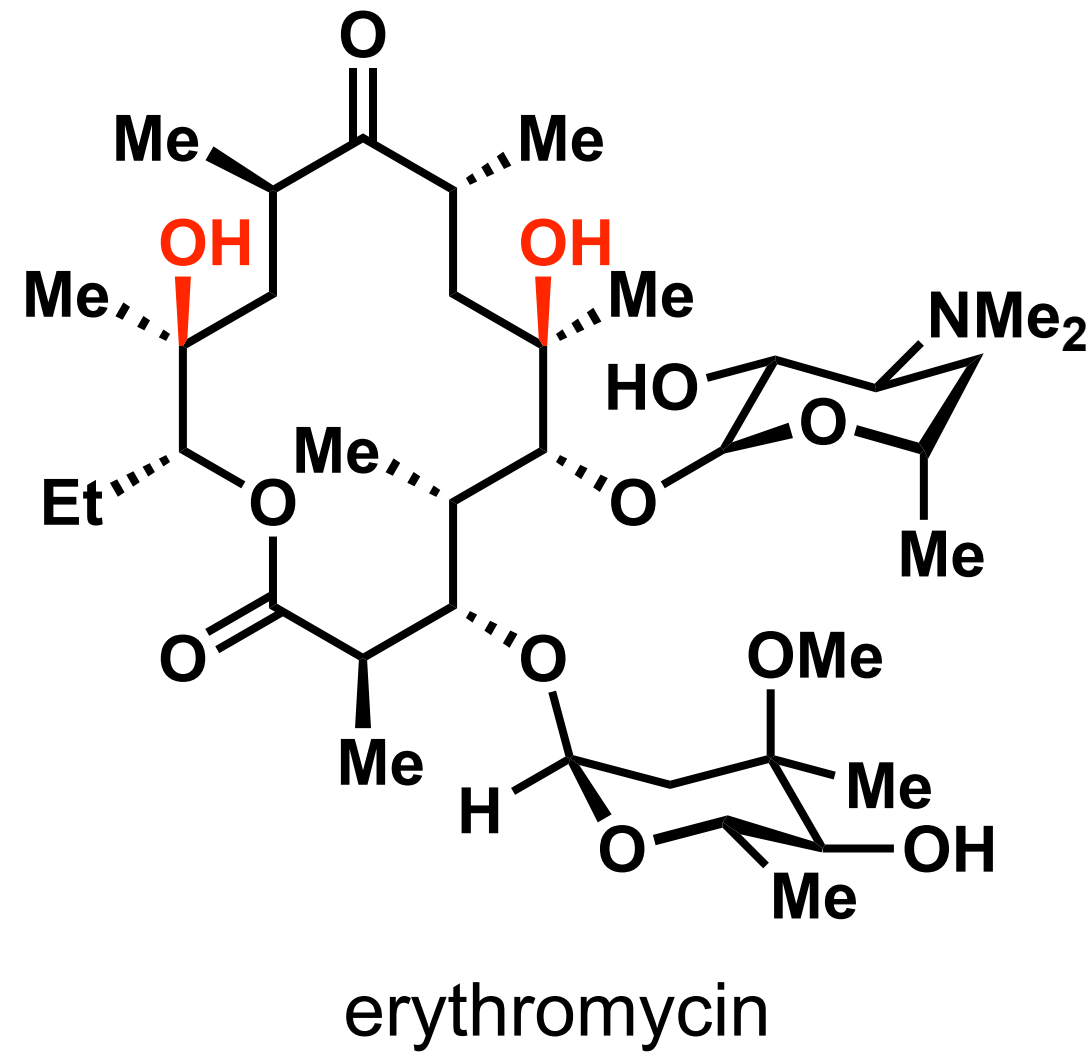
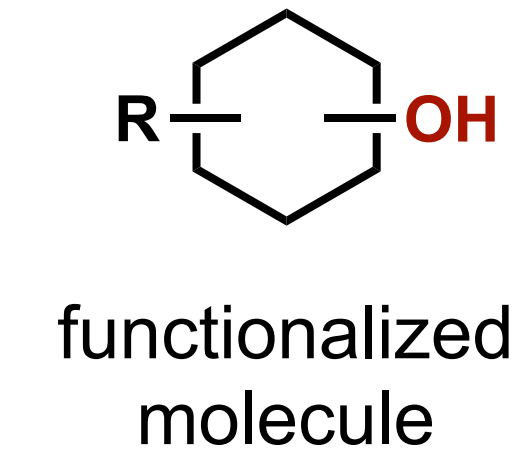
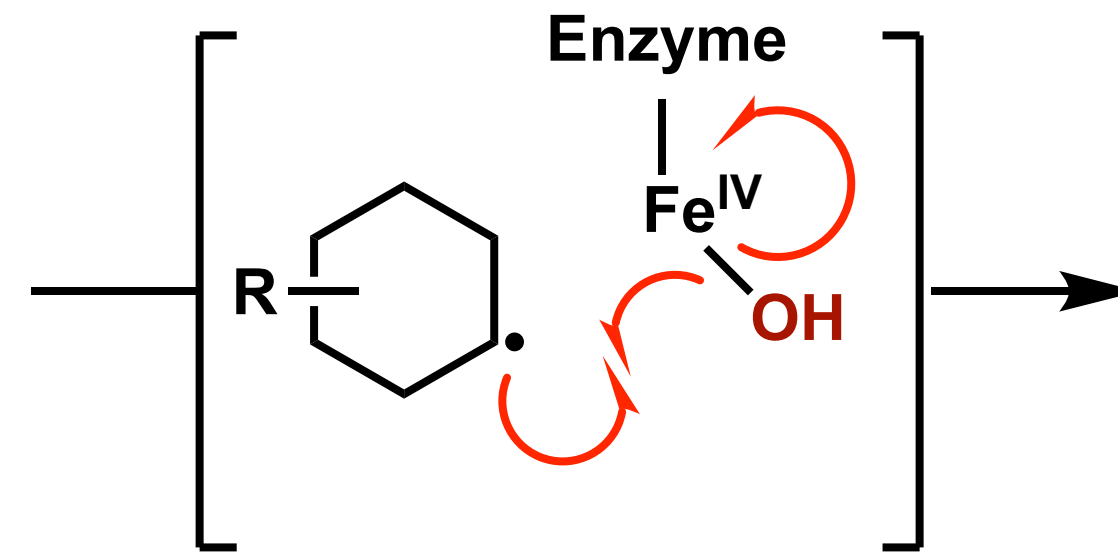
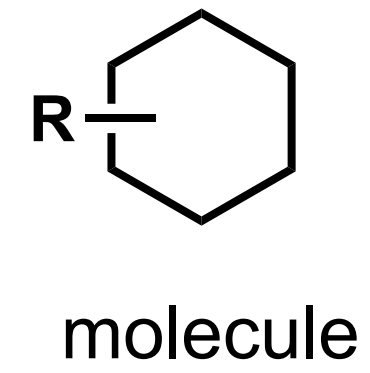
common anti-inflammatory



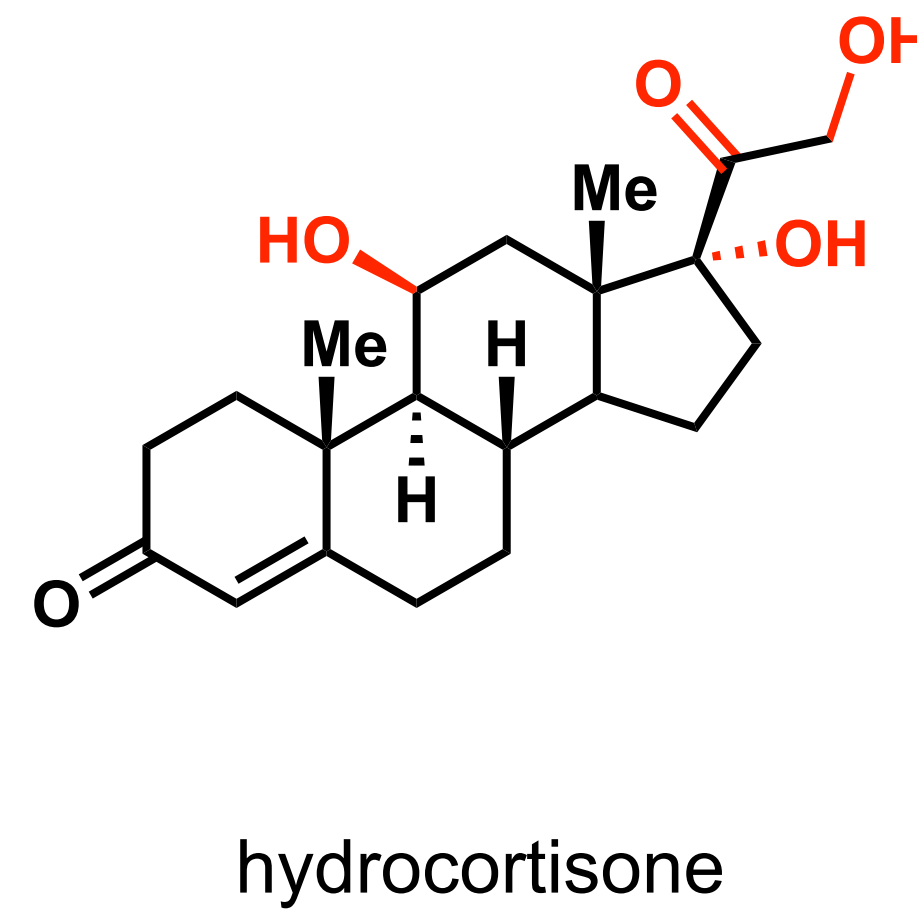
Pravachol

high cholesterol treatment

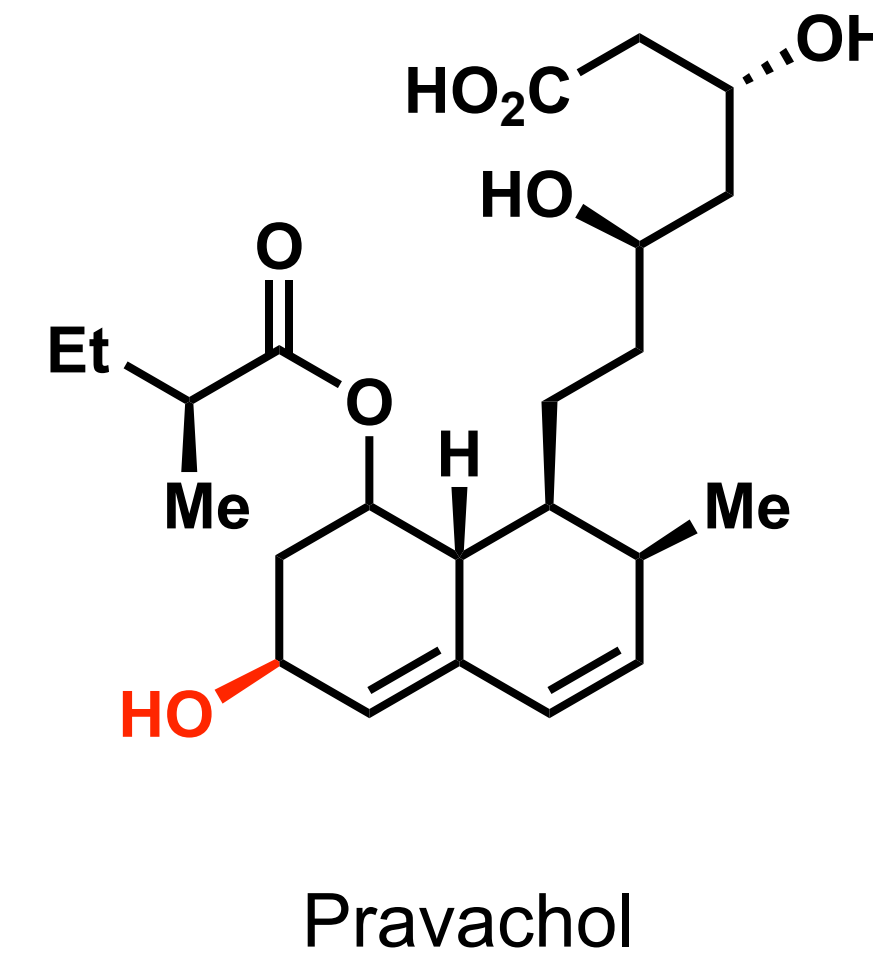
A Different Reaction Altogether



common anti-bacterial

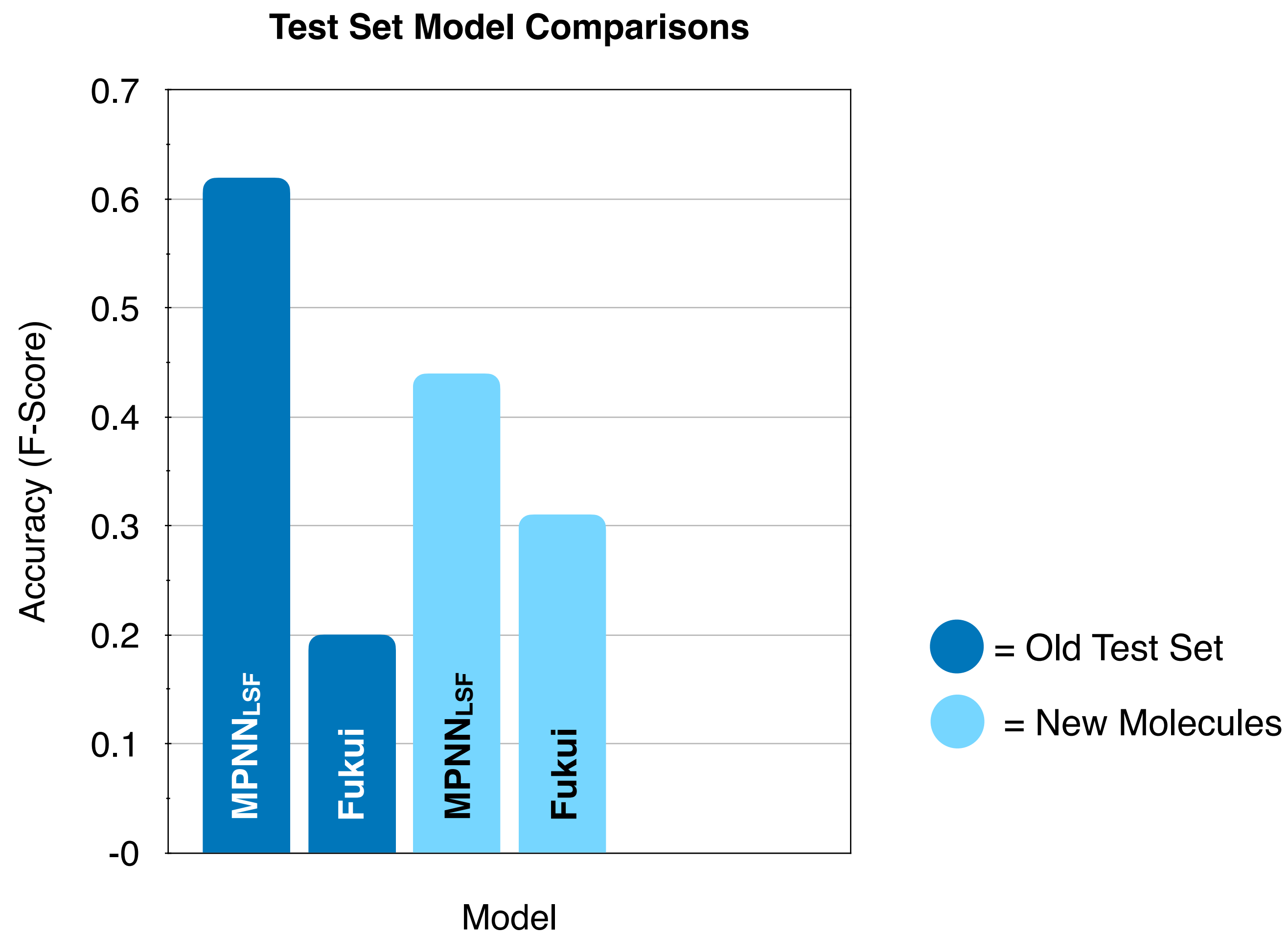


common anti-inflammatory

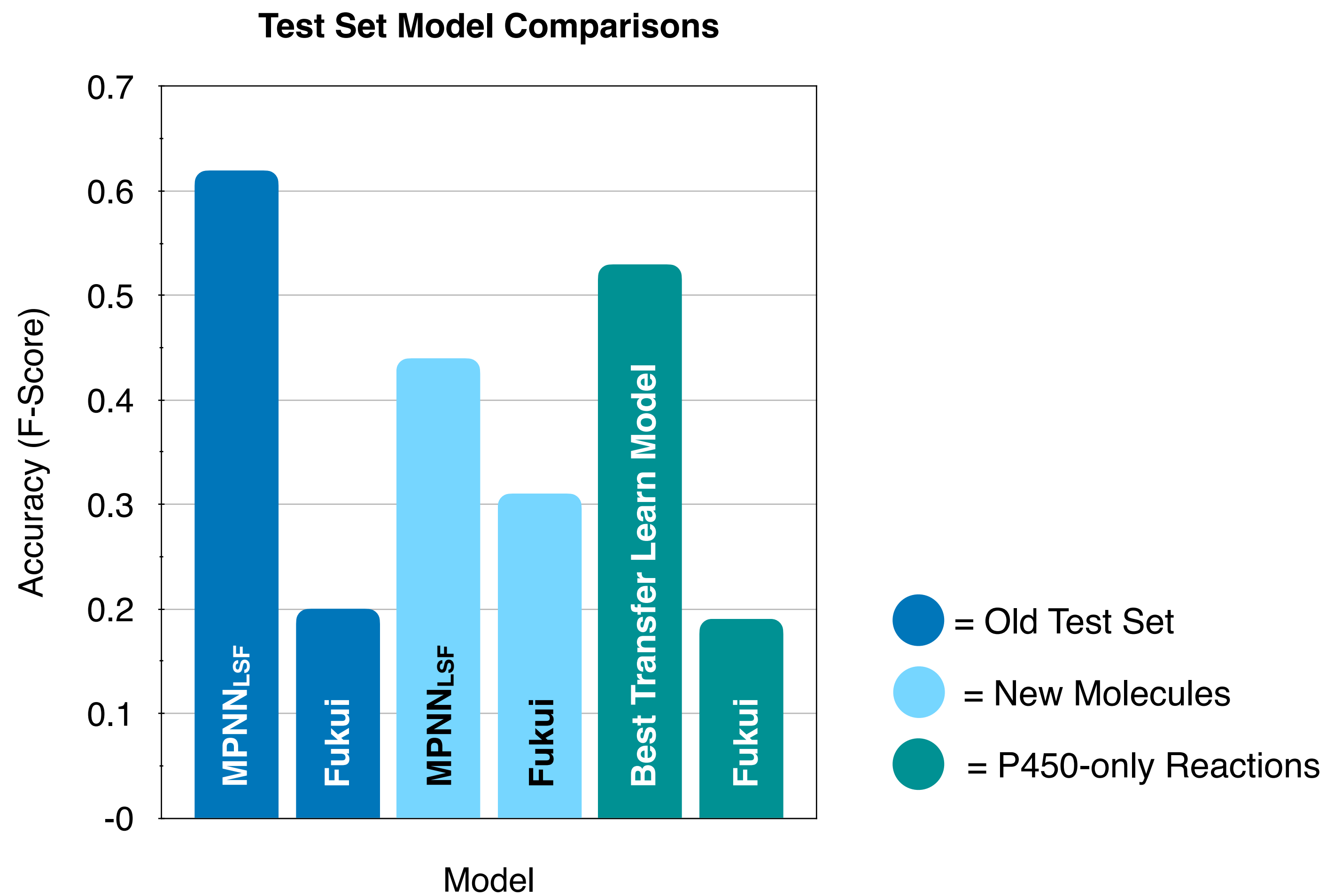


high cholesterol treatment

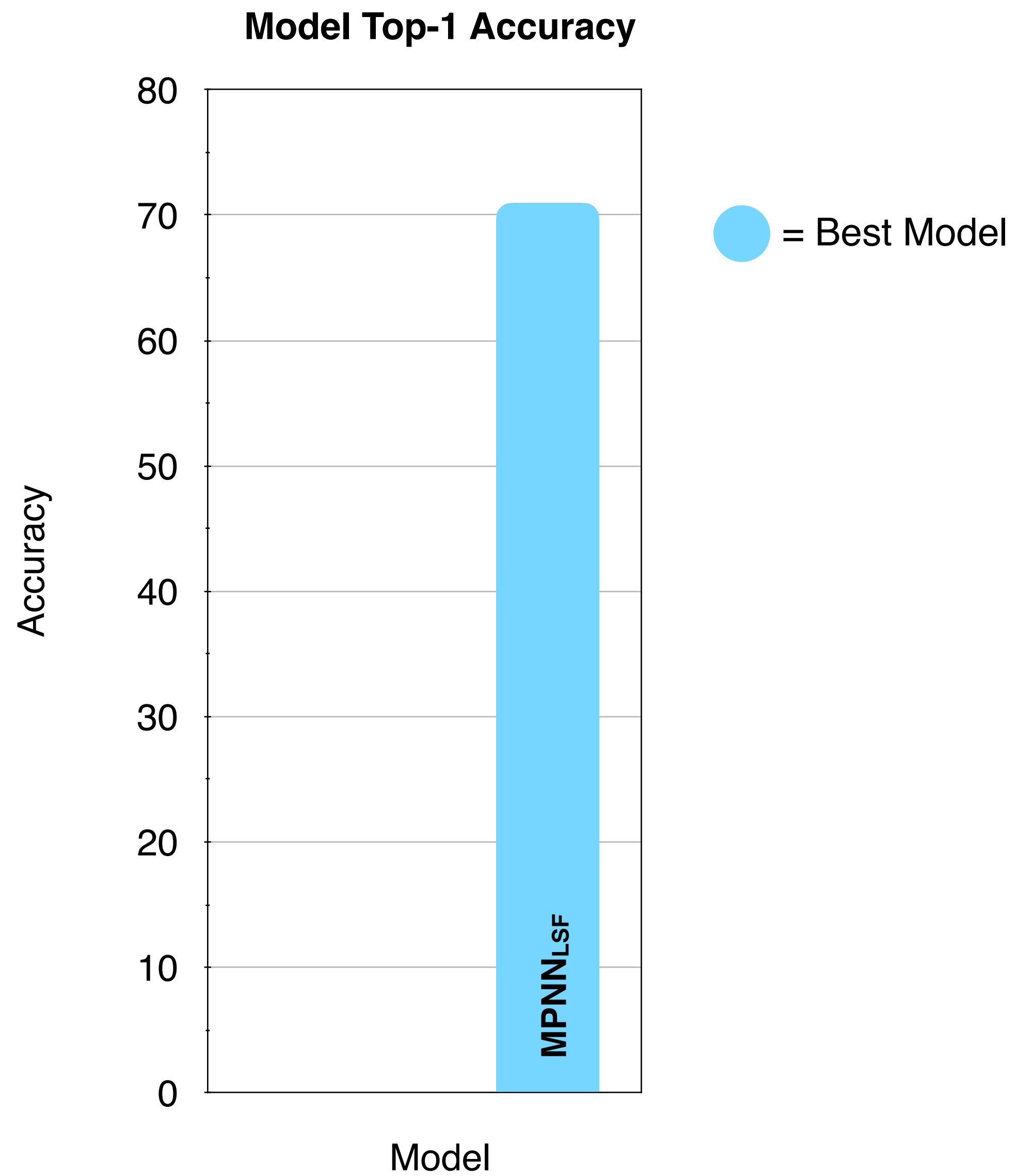
How Do We Perform on Completely Unseen Molecules?



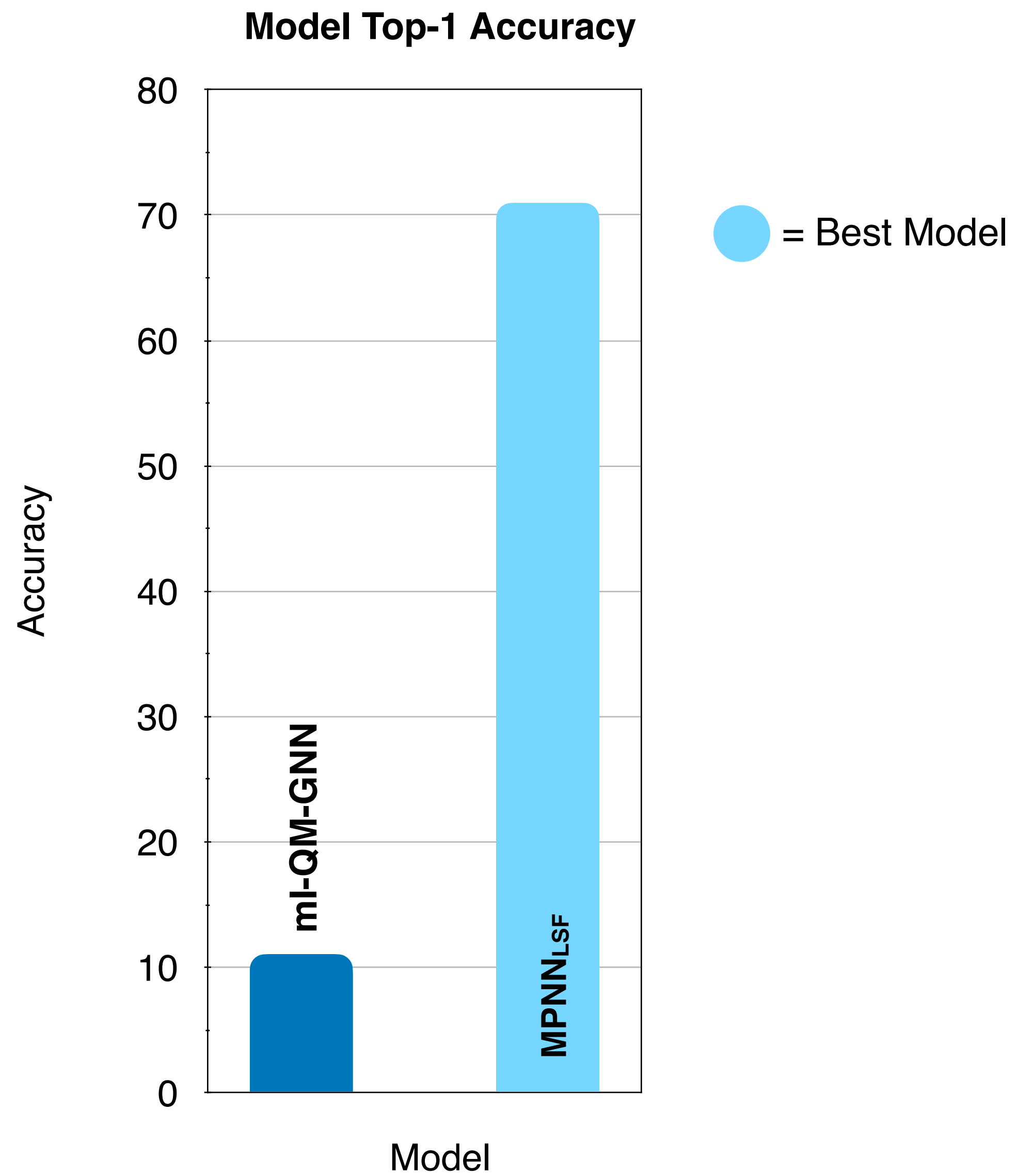
How Do We Perform On a P450-Only Test Set?



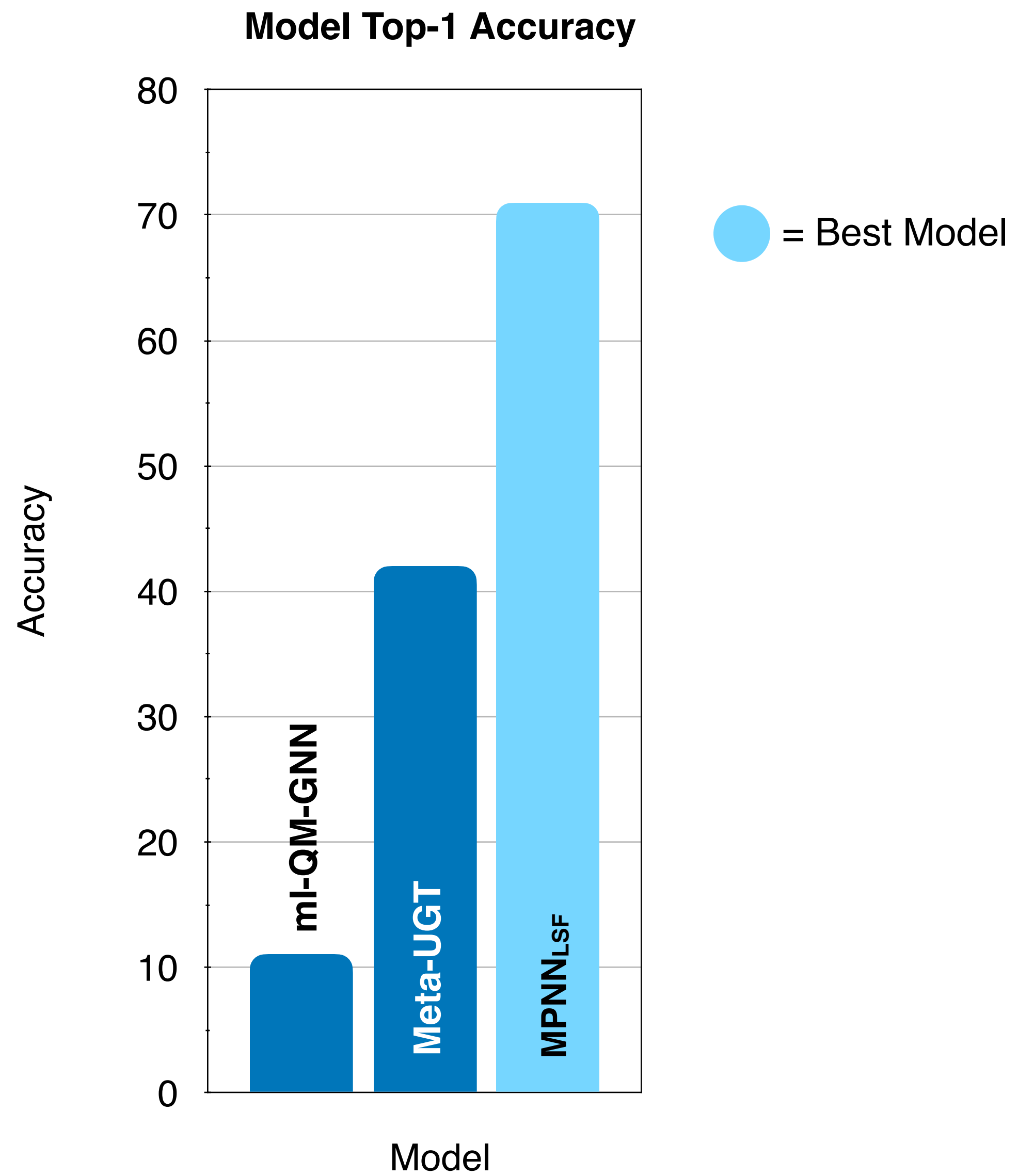
Comparison To Other Reactivity-Based Models

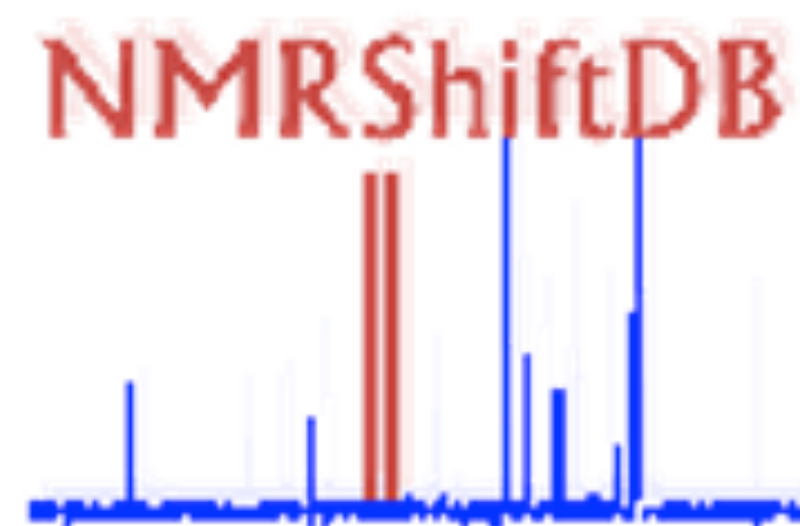


Comparison To Other Reactivity-Based Models



Comparison To Other Reactivity-Based Models

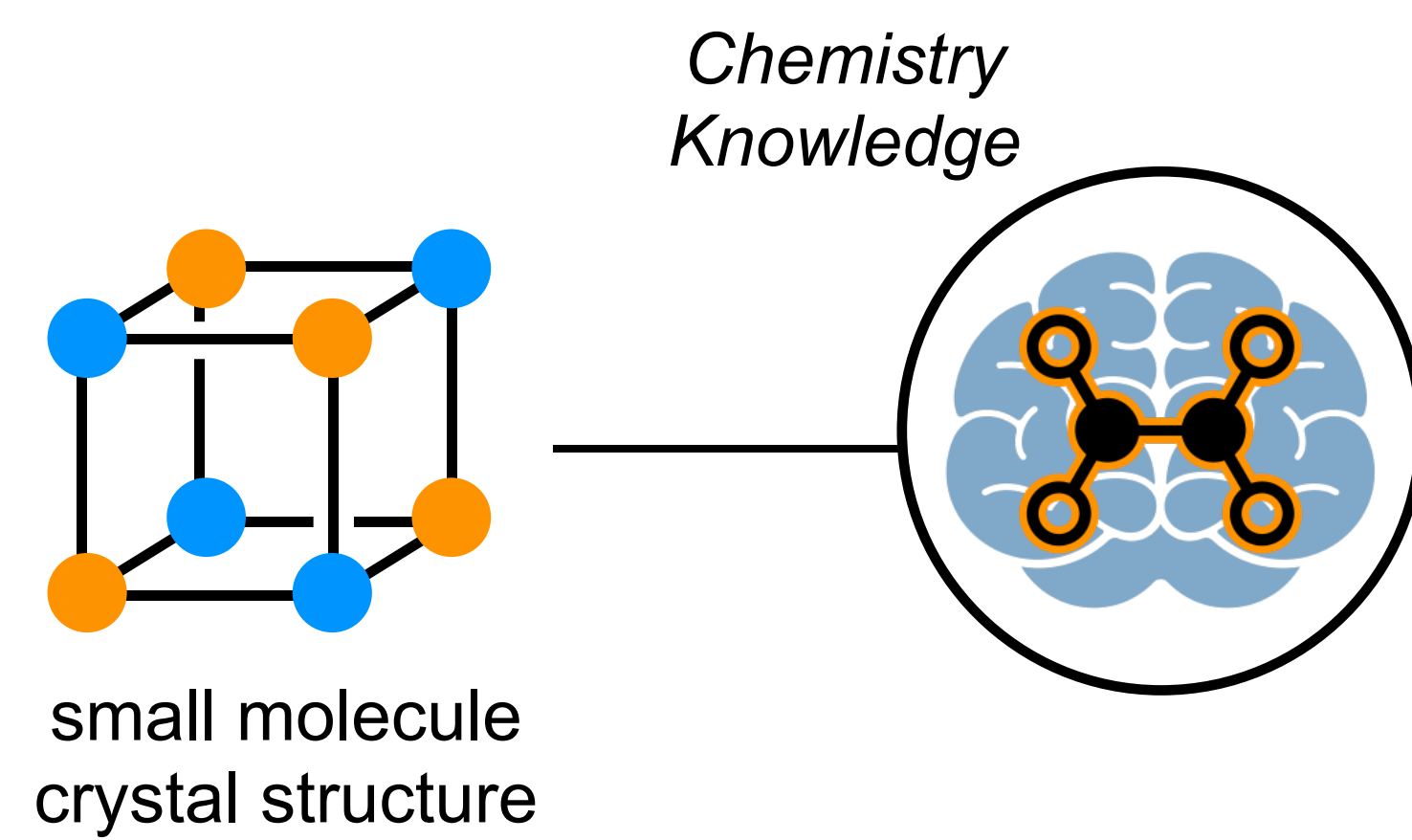


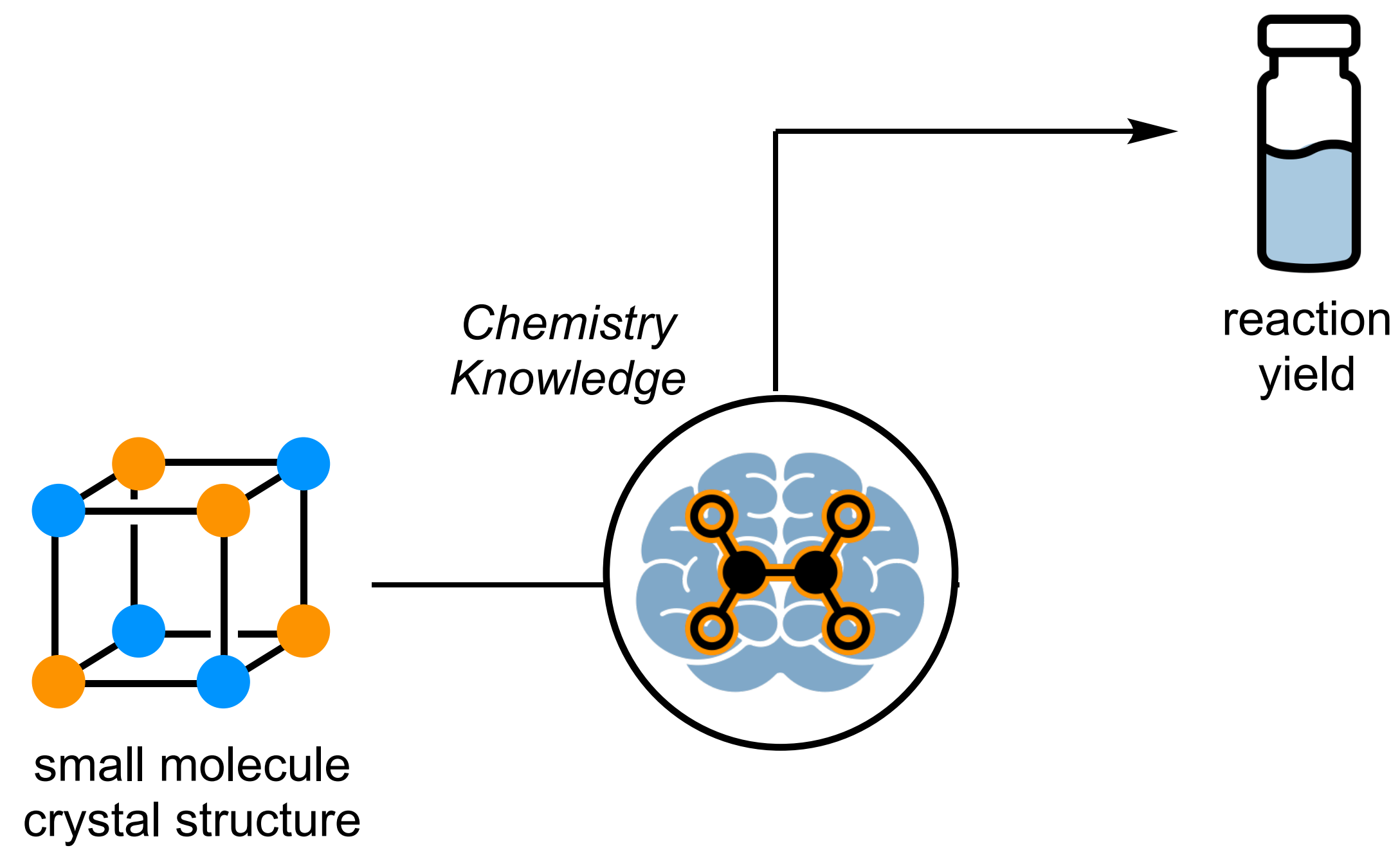


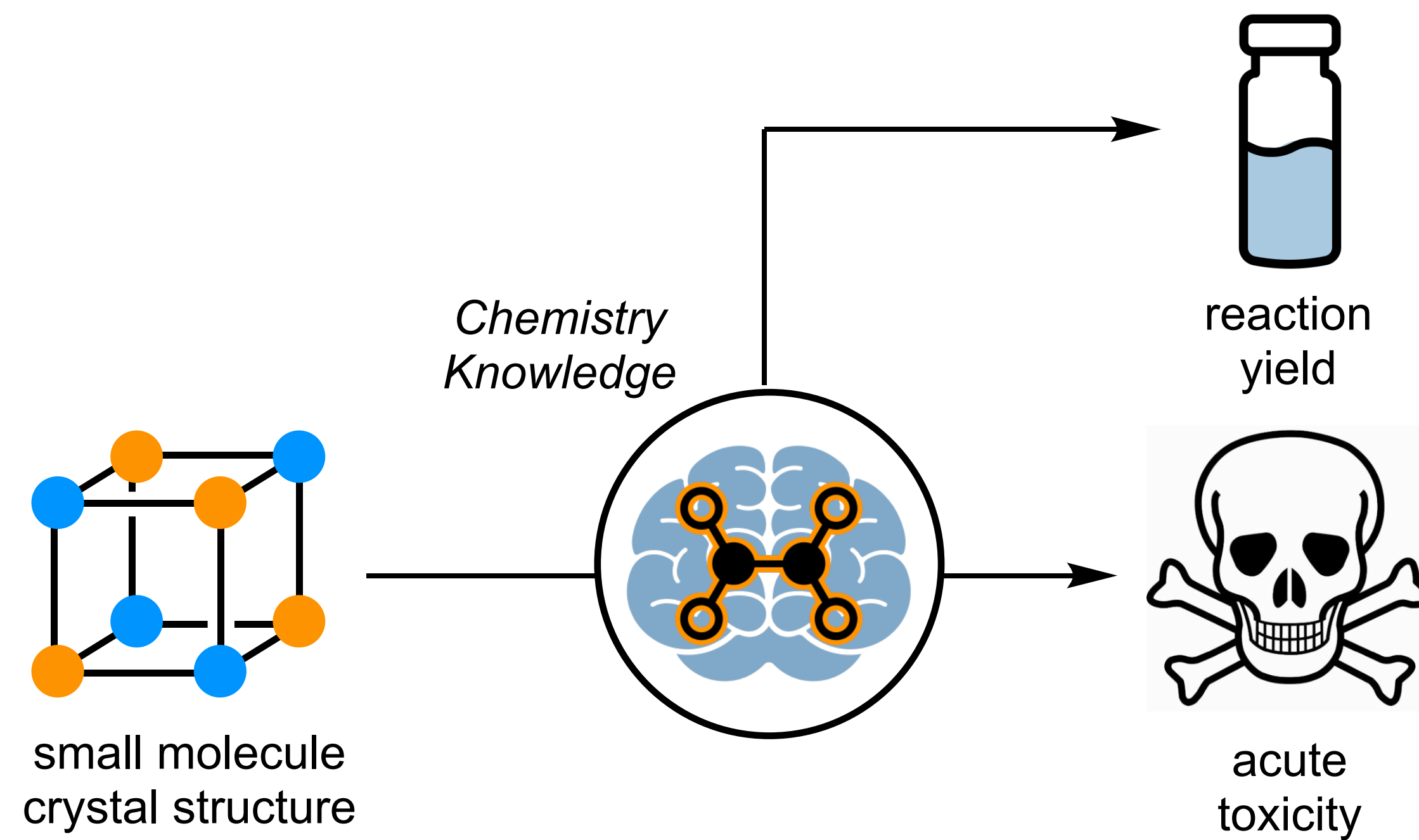
~27,000 spectra

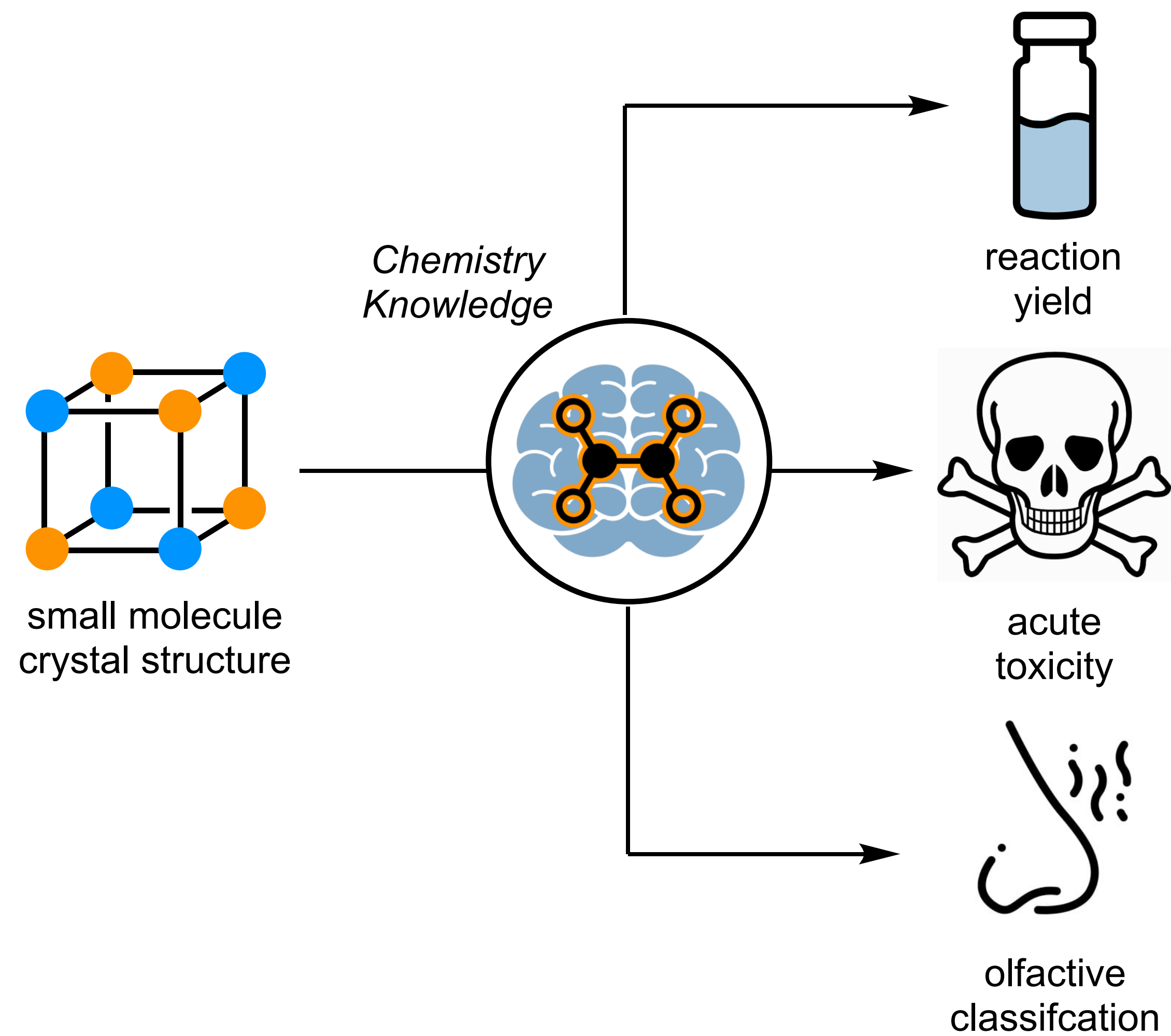
The logo for the Cambridge Crystallographic Data Centre (CCDC), featuring the letters 'CCDC' in a bold, black, sans-serif font. A small blue dot is positioned below the 'D'.

1,000,000+ compounds









Suzuki and Buchwald-Hartwig Cross Coupling Yield Predictions

Model	Suzuki Yield Error (MAE)		Buchwald-Hartwig Yield Error (MAE)			
	Unseen Boronic Acids	Unseen Aryl Halides	Unseen Boronic Acids	Unseen Aryl Halides	Unseen Ligands	Unseen Additives
Random Forest						
Adaboost						
Yield-BERT						
GraphRXN						
Crystal-Yield						

*Increased Crystal-Yield's size to half of GraphRXN's parameters

Suzuki and Buchwald-Hartwig Cross Coupling Yield Predictions

Model	Suzuki Yield Error (MAE)		Buchwald-Hartwig Yield Error (MAE)			
	Unseen Boronic Acids	Unseen Aryl Halides	Unseen Boronic Acids	Unseen Aryl Halides	Unseen Ligands	Unseen Additives
Random Forest						
Adaboost						
Yield-BERT						
GraphRXN						
Crystal-Yield	18.4 ± 0.3	18.5 ± 0.2	21.3 ± 3.3	13.4 ± 0.3	11.7 ± 2.2*	16.2 ± 0.4

*Increased Crystal-Yield's size to half of GraphRXN's parameters

Model	Suzuki Yield Error (MAE)		Buchwald-Hartwig Yield Error (MAE)			
	Unseen Boronic Acids	Unseen Aryl Halides	Unseen Boronic Acids	Unseen Aryl Halides	Unseen Ligands	Unseen Additives
Random Forest	19.5 ± 0.03	19.5 ± 0.03	25.2 ± 2.0	28.1 ± 4.1	28.5 ± 0.6	30.4 ± 1.5
Adaboost	21.6 ± 0.1	21.5 ± 0.1	24.7 ± 2.6	25.5 ± 2.9	27.9 ± 0.7	26.7 ± 0.5
Yield-BERT						
GraphRXN						
Crystal-Yield	18.4 ± 0.3	18.5 ± 0.2	21.3 ± 3.3	13.4 ± 0.3	11.7 ± 2.2*	16.2 ± 0.4

*Increased Crystal-Yield's size to half of GraphRXN's parameters

Model	Suzuki Yield Error (MAE)		Buchwald-Hartwig Yield Error (MAE)			
	Unseen Boronic Acids	Unseen Aryl Halides	Unseen Boronic Acids	Unseen Aryl Halides	Unseen Ligands	Unseen Additives
Random Forest	19.5 ± 0.03	19.5 ± 0.03	25.2 ± 2.0	28.1 ± 4.1	28.5 ± 0.6	30.4 ± 1.5
Adaboost	21.6 ± 0.1	21.5 ± 0.1	24.7 ± 2.6	25.5 ± 2.9	27.9 ± 0.7	26.7 ± 0.5
Yield-BERT	21.9 ± 0.06	22.0 ± 0.03	24.7 ± 2.1	24.3 ± 1.6	24.3 ± 1.4	24.1 ± 0.7
GraphRXN	40.0 ± 3.0	37.8 ± 2.7	25.2 ± 7.0	17.9 ± 4.6	13.8 ± 1.7	17.5 ± 1.8
Crystal-Yield	18.4 ± 0.3	18.5 ± 0.2	21.3 ± 3.3	13.4 ± 0.3	11.7 ± 2.2*	16.2 ± 0.4

*Increased Crystal-Yield's size to half of GraphRXN's parameters

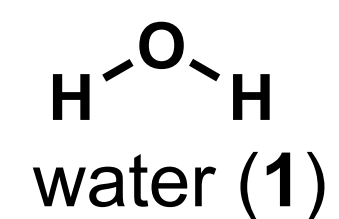
Model	Pharmaceuticals (MAE)
Random Forest	
Gaussian Process	
Adaboost	
Oloren Chem Engine	
Crystal-Tox	0.52 ± 0.007

Model	Pharmaceuticals (MAE)
Random Forest	0.62 ± 0.002
Gaussian Process	0.73 ± 0.002
Adaboost	0.71 ± 0.002
Oloren Chem Engine	
Crystal-Tox	0.52 ± 0.007

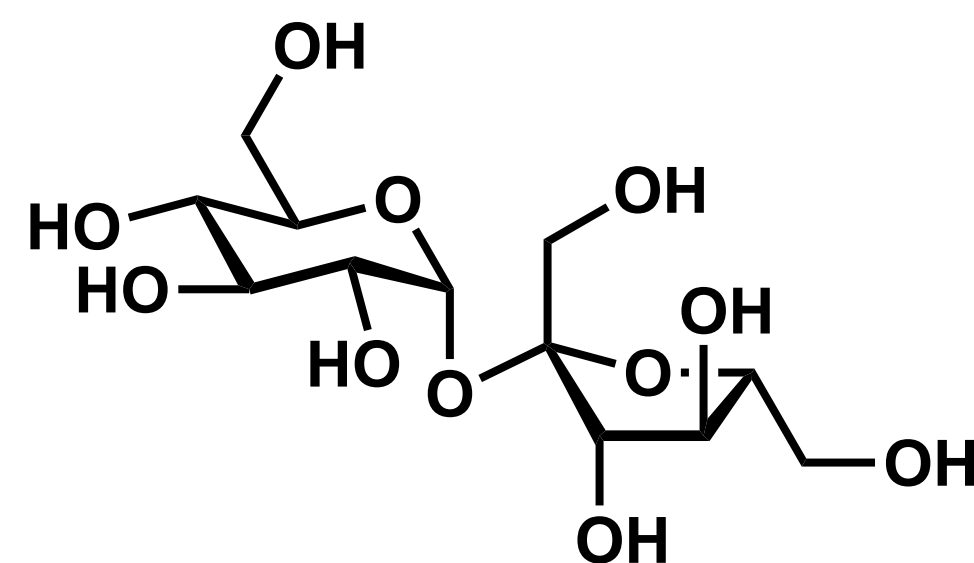
Model	Pharmaceuticals (MAE)
Random Forest	0.62 ± 0.002
Gaussian Process	0.73 ± 0.002
Adaboost	0.71 ± 0.002
Oloren Chem Engine	0.55 ± 0.009
Crystal-Tox	0.52 ± 0.007

New Molecules for Testing

Benign Molecules

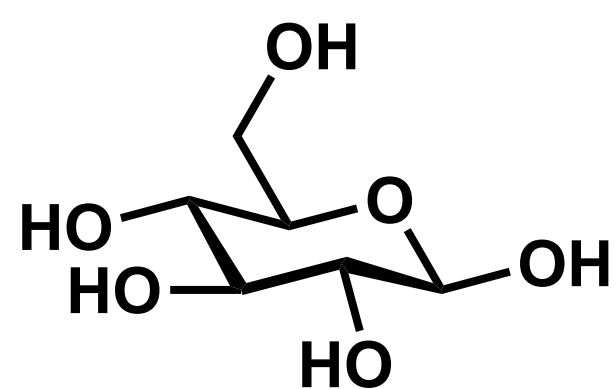


LD50 90,000



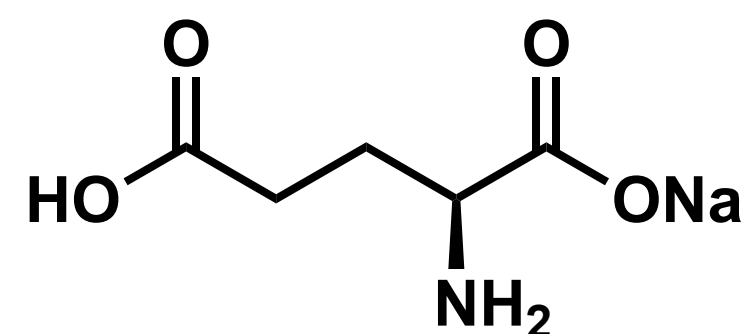
sucrose (2)

LD50 29,700



glucose (3)

LD50 = 25,800

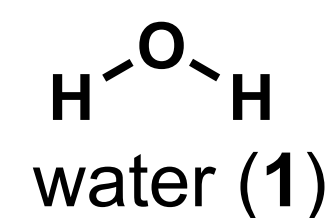


monosodium glutamate (4)

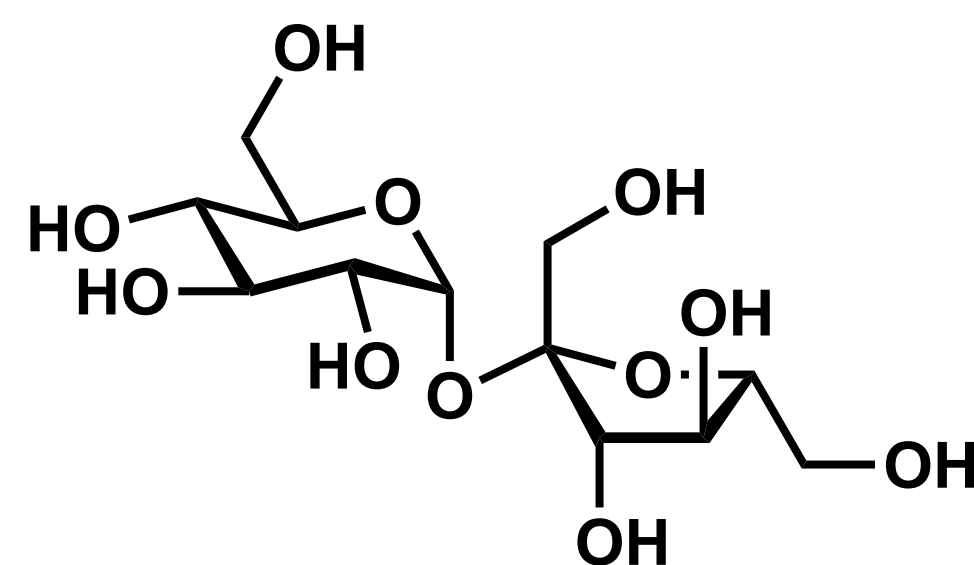
LD50 = 16,600

Benign Molecules

Natural Toxins

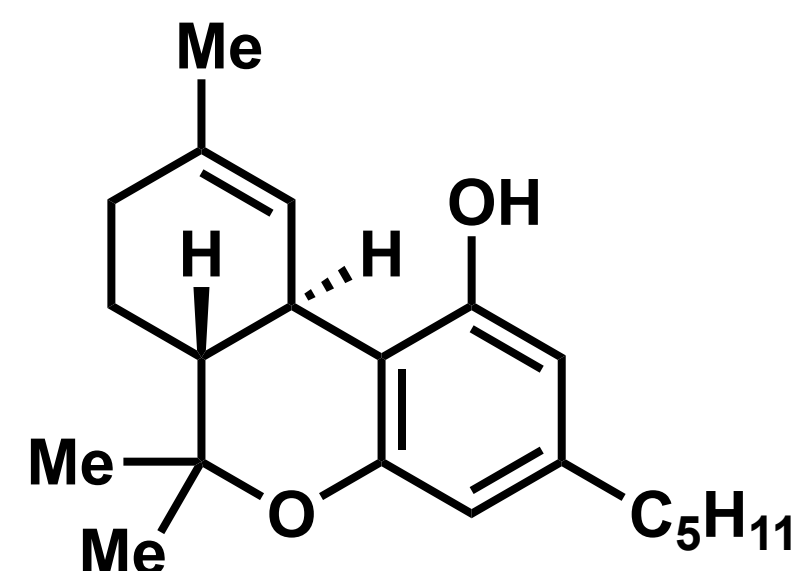


LD50 90,000



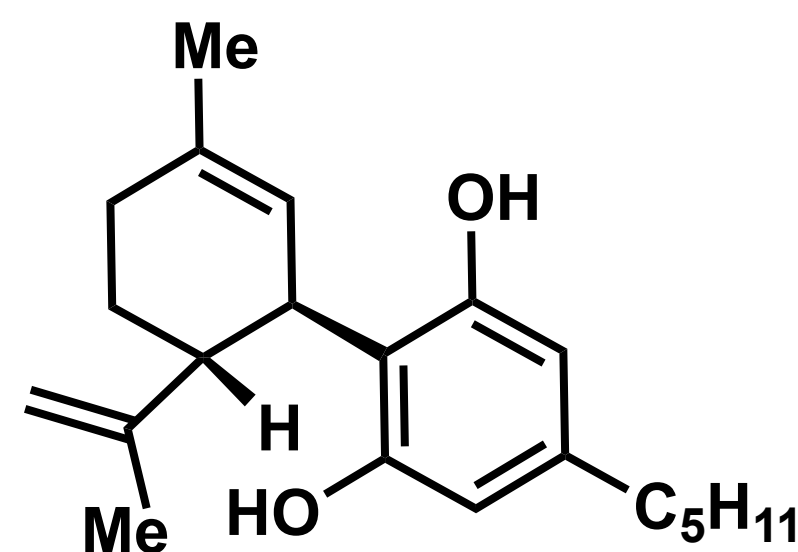
sucrose (2)

LD50 29,700



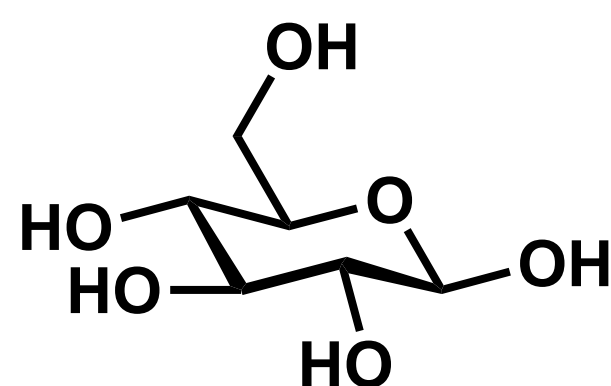
THC (5)

LD50 = 1,270



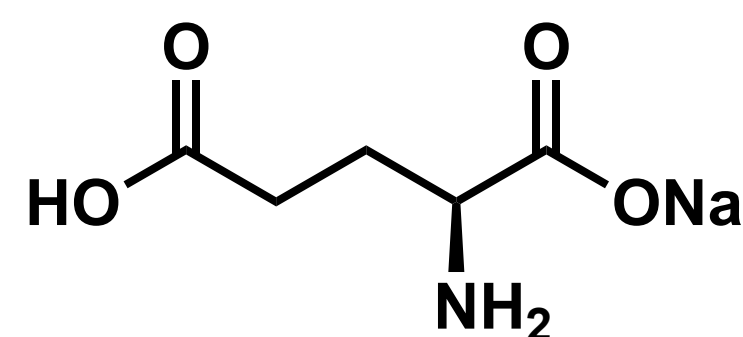
CBD (6)

LD50 = 980



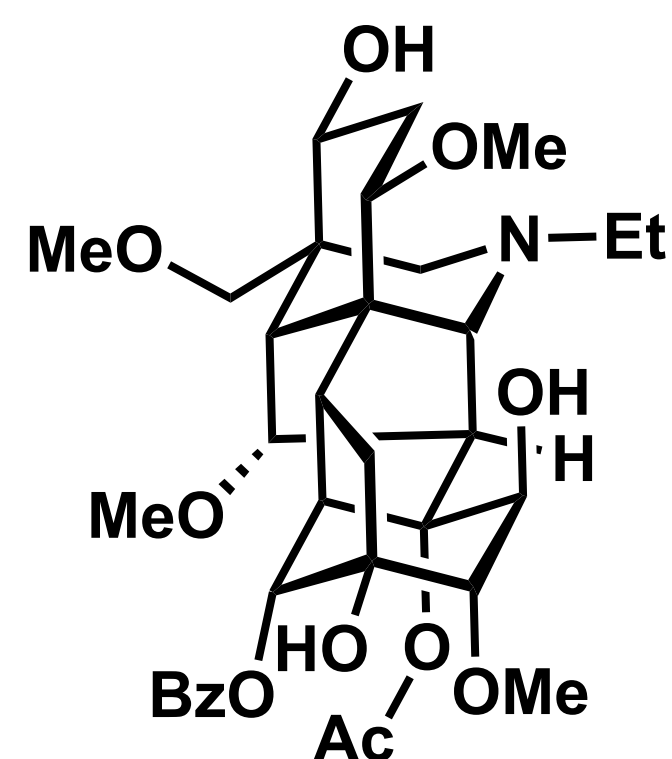
glucose (3)

LD50 = 25,800



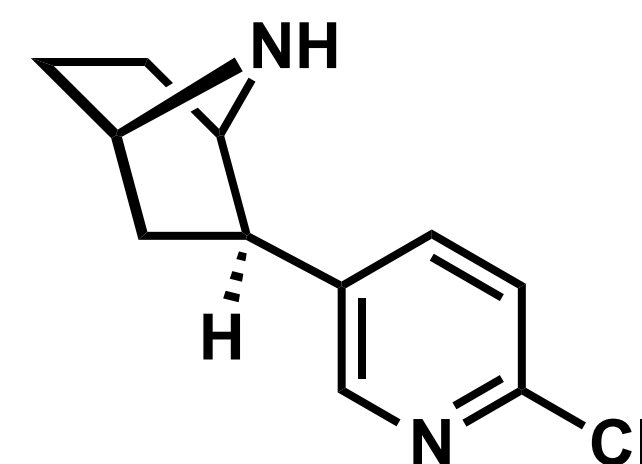
monosodium glutamate (4)

LD50 = 16,600



aconitine (7)

LD50 = 0.08



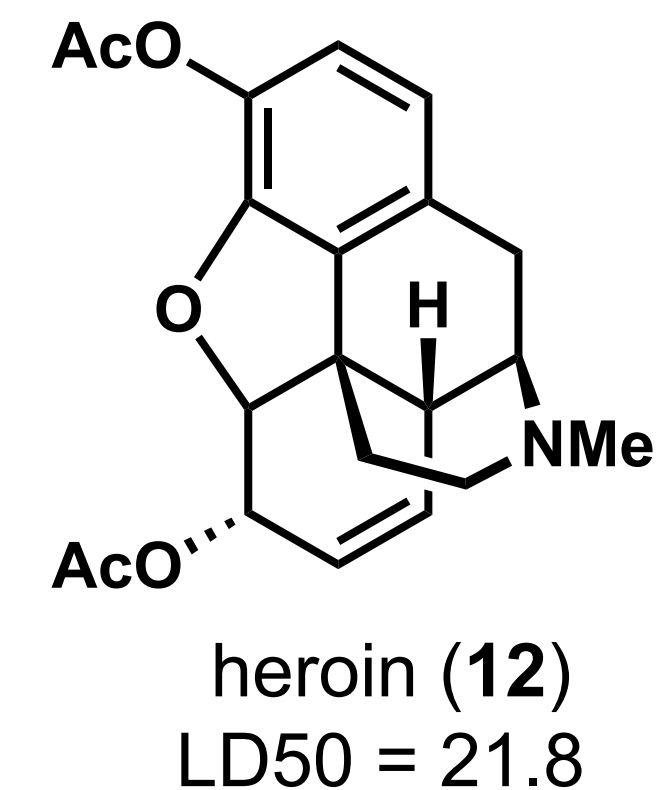
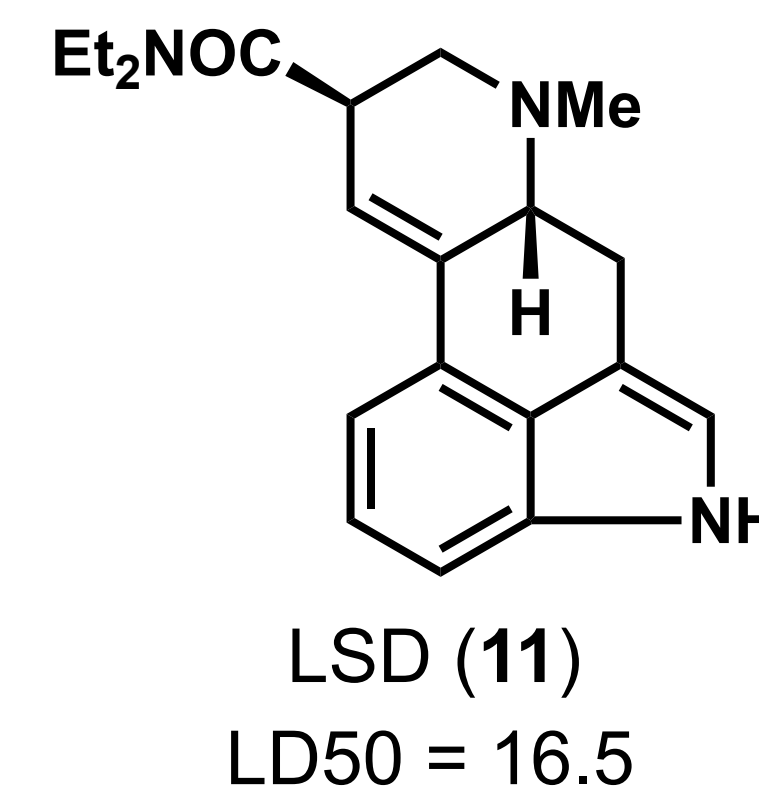
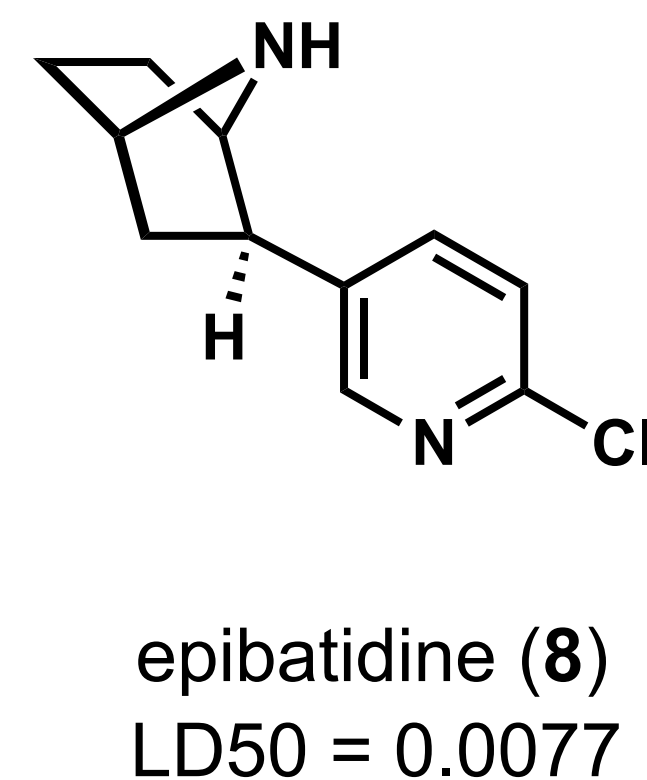
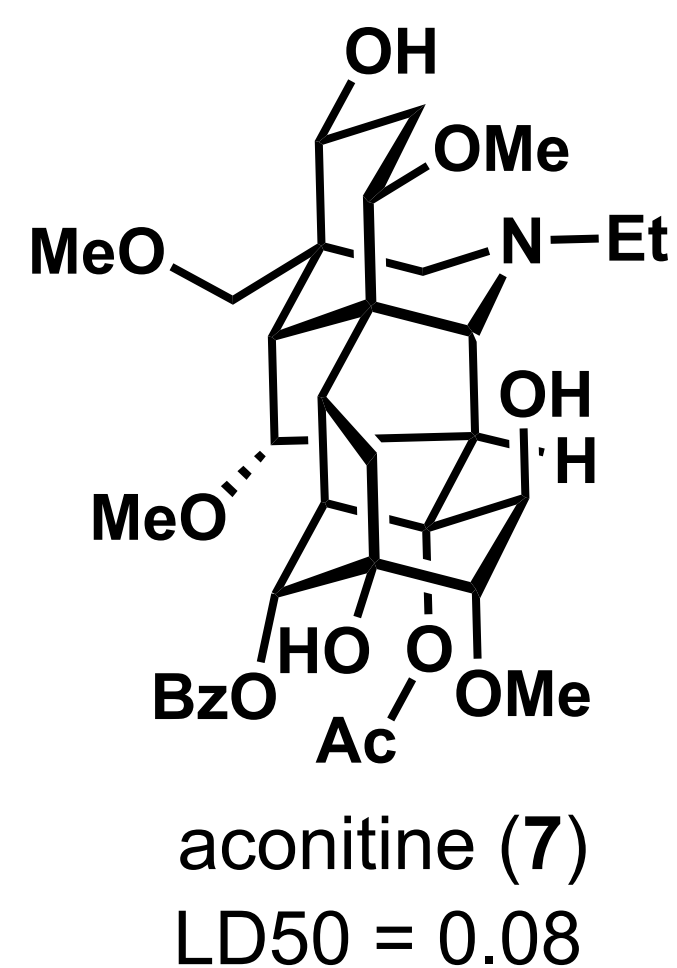
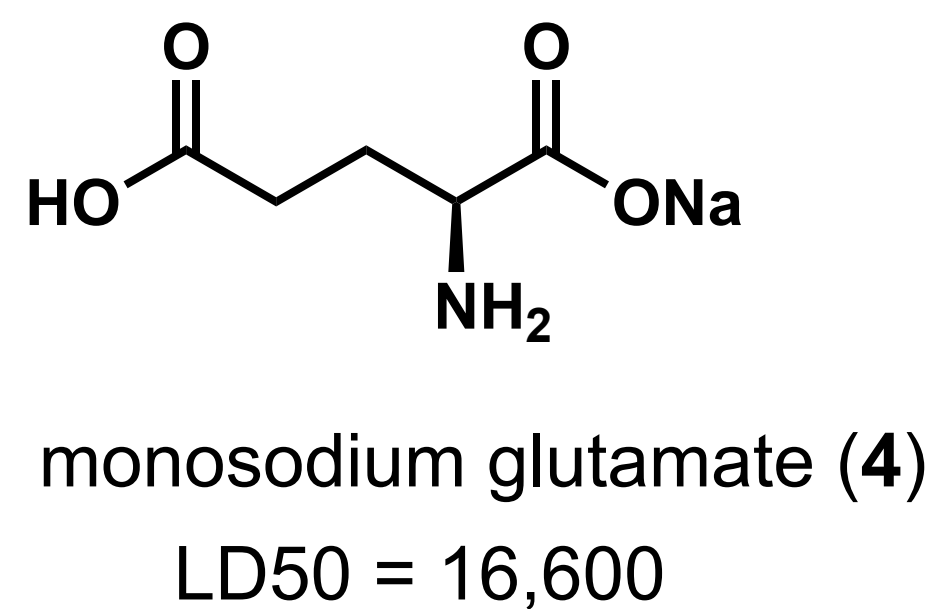
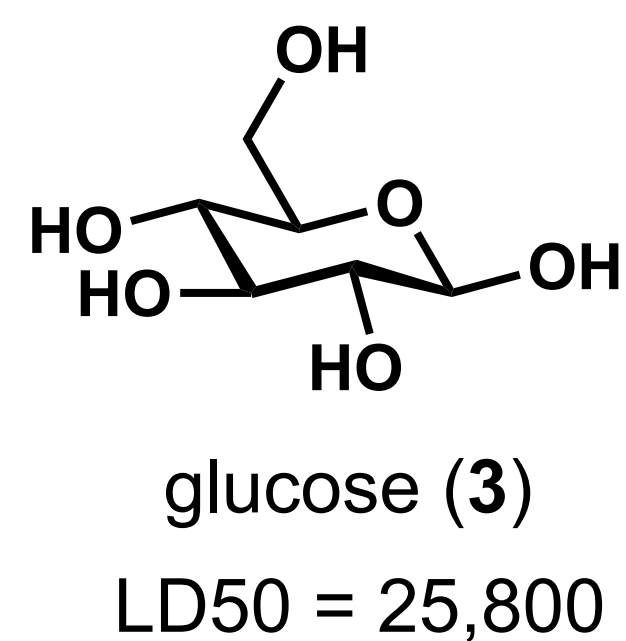
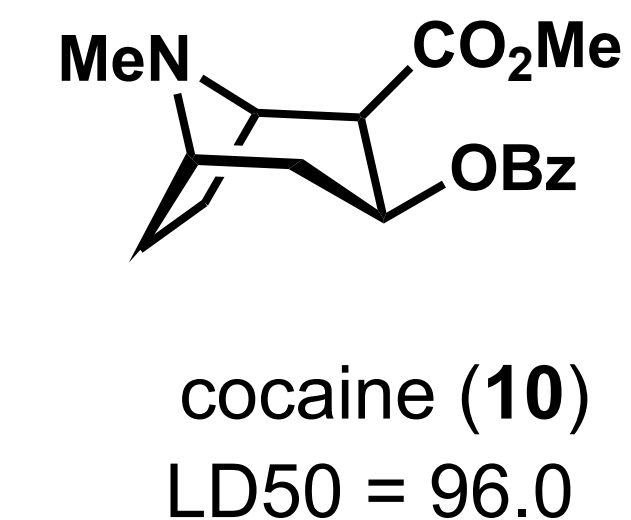
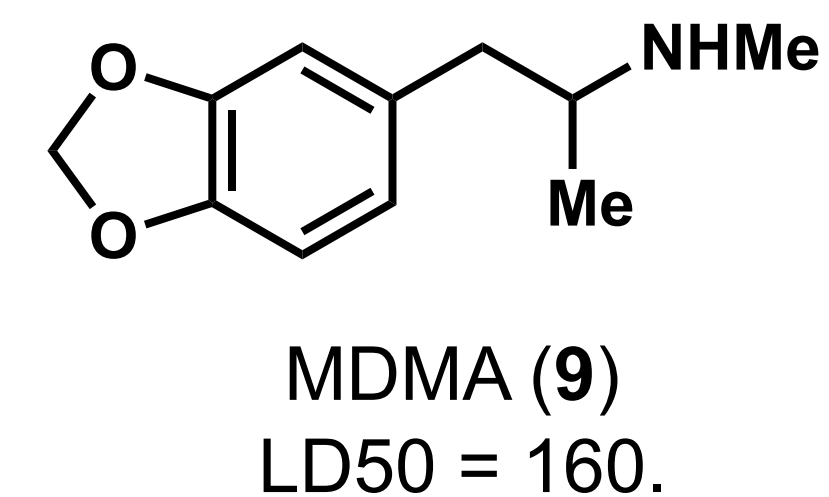
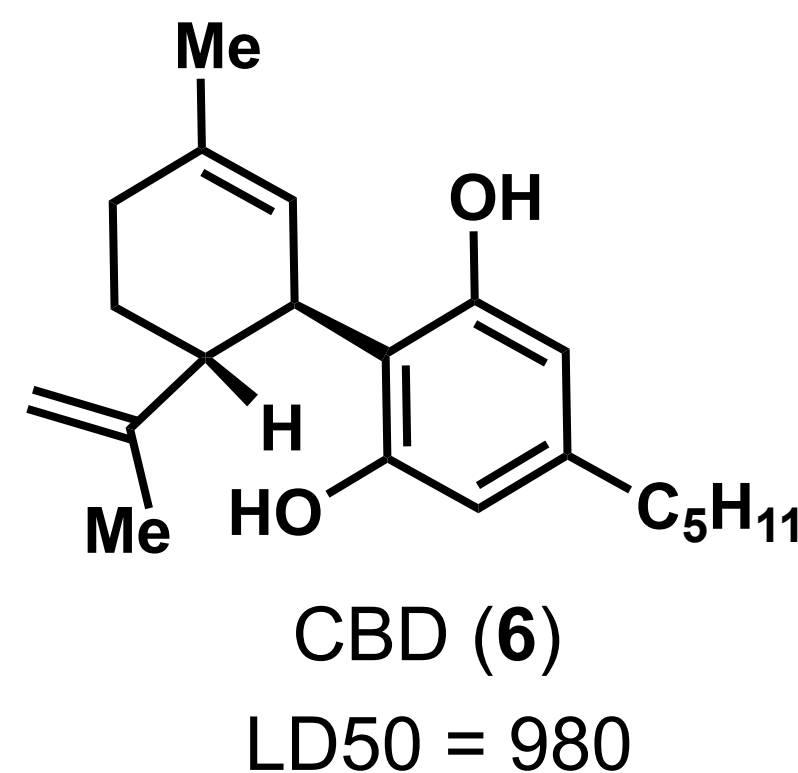
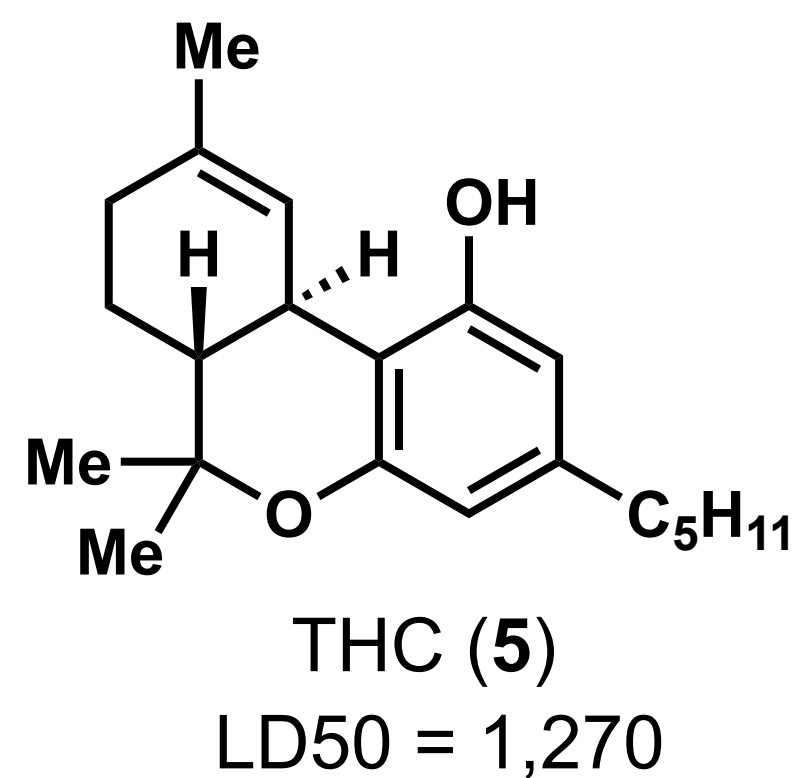
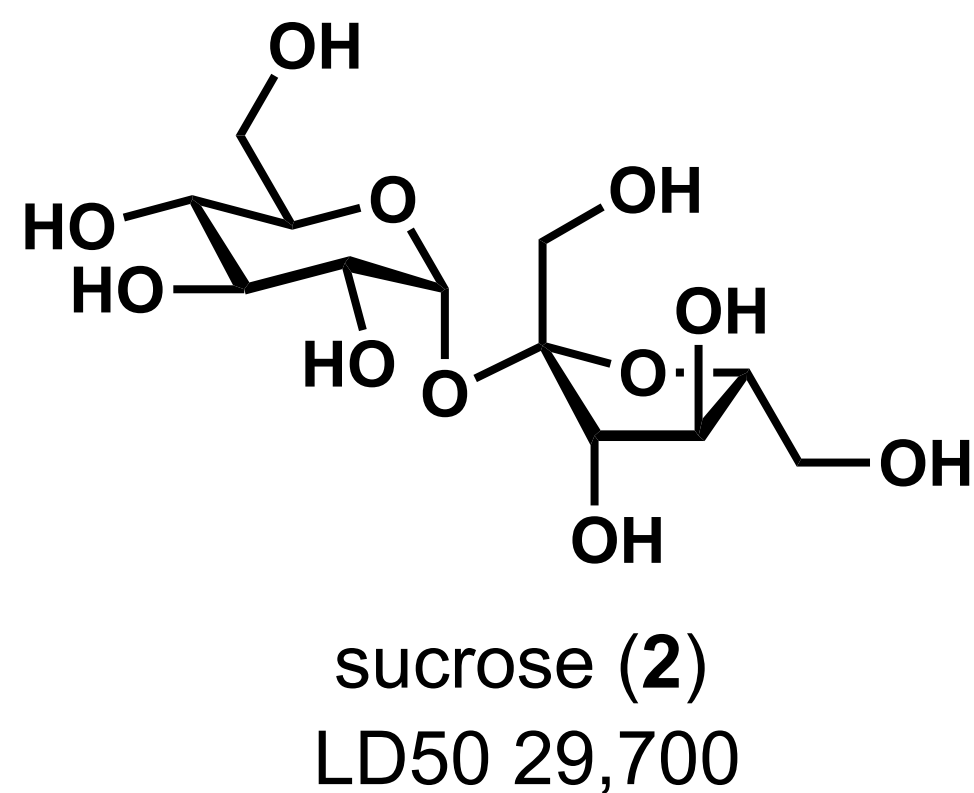
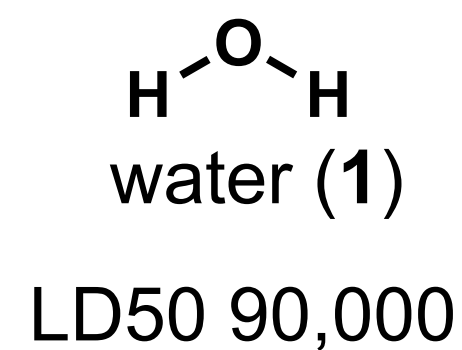
epibatidine (8)

LD50 = 0.0077

Benign Molecules

Natural Toxins

Illicit Substances



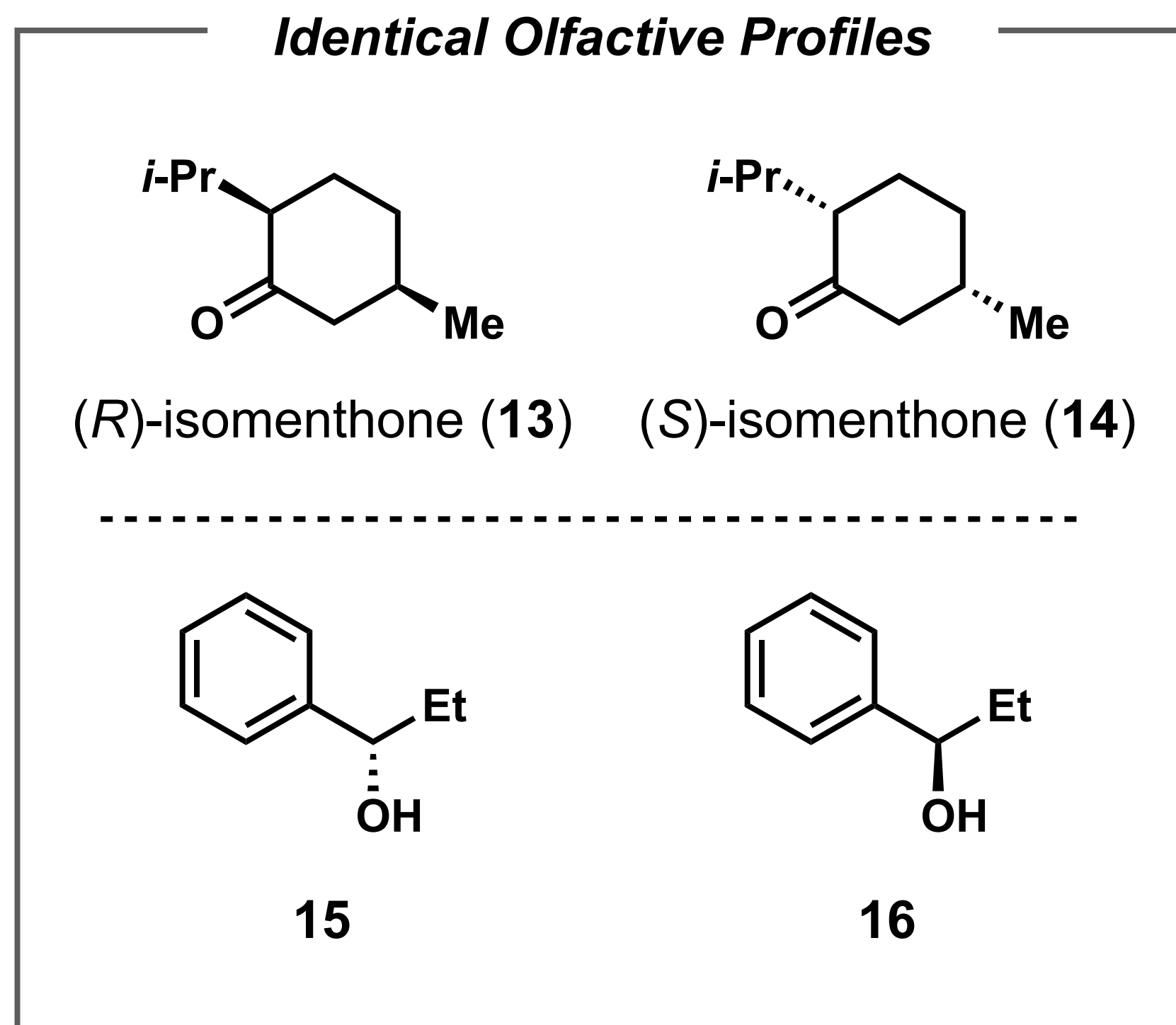
Model	Pharmaceuticals (MAE)
Random Forest	0.62 ± 0.002
Gaussian Process	0.73 ± 0.002
Adaboost	0.71 ± 0.002
Oloren Chem Engine	0.55 ± 0.009
Crystal-Tox	0.52 ± 0.007

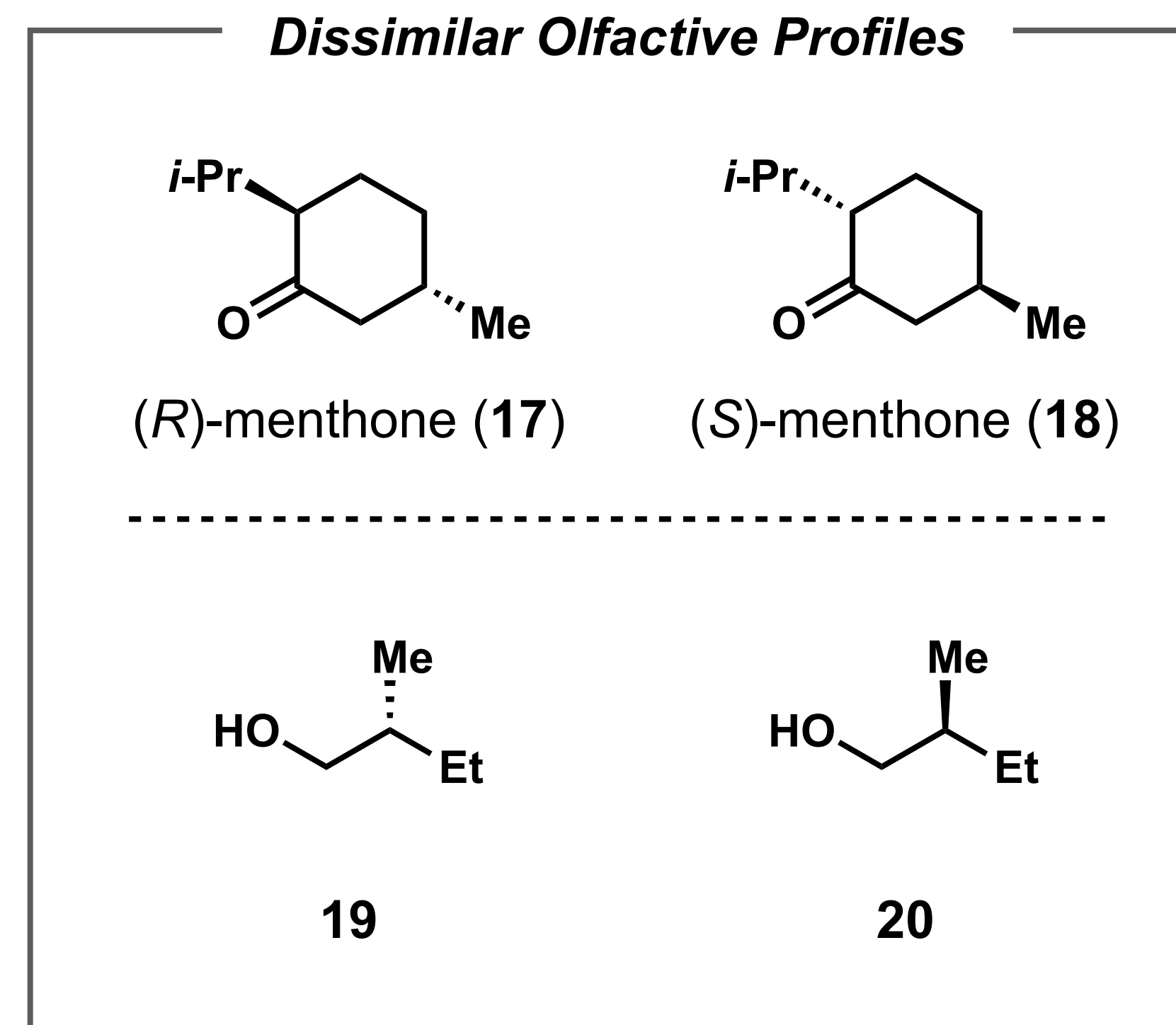
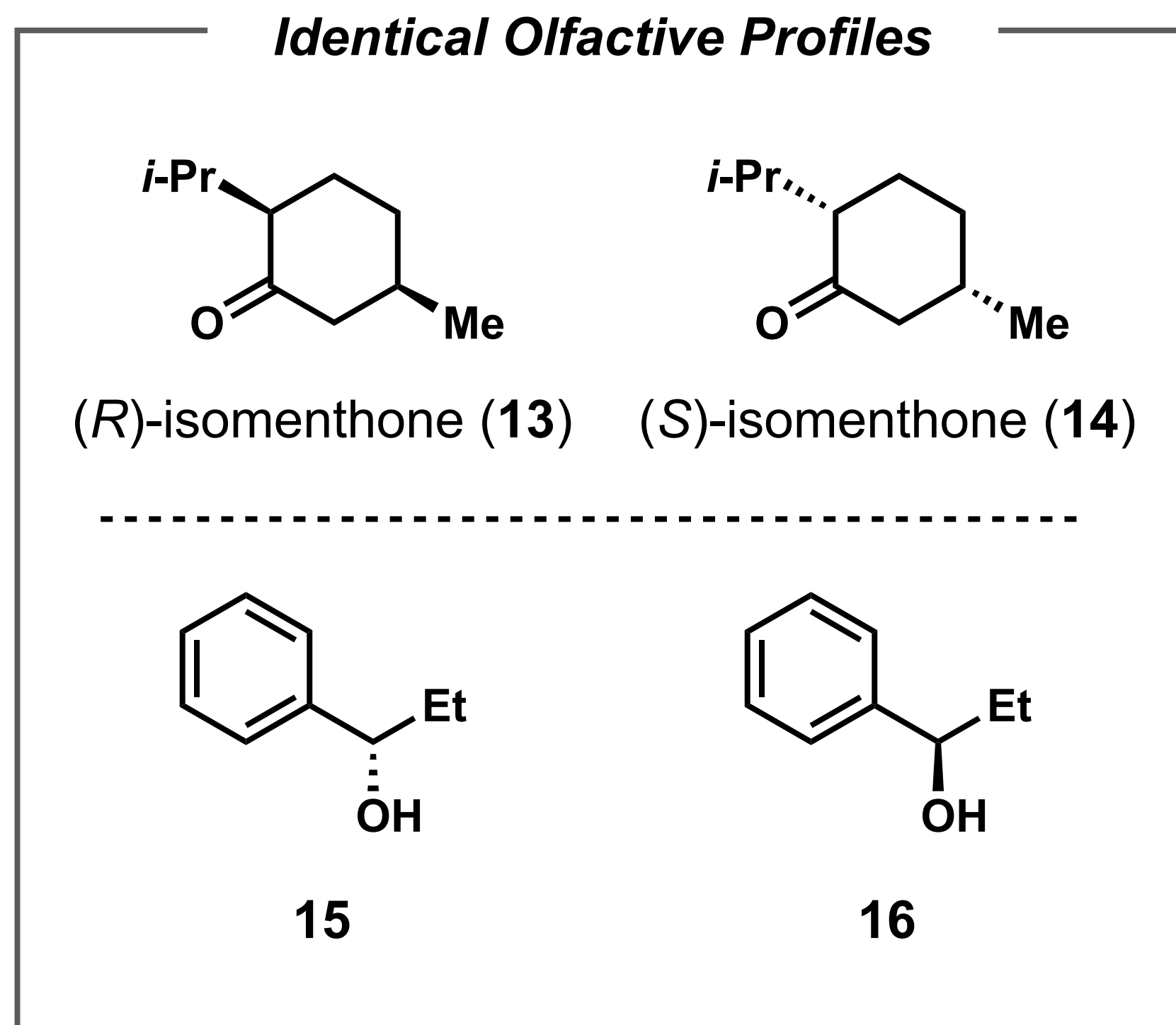
Model	Pharmaceuticals (MAE)	Non-Pharmaceuticals (MAE)
Random Forest	0.62 ± 0.002	1.59 ± 0.02
Gaussian Process	0.73 ± 0.002	1.86 ± 0.002
Adaboost	0.71 ± 0.002	1.77 ± 0.002
Oloren Chem Engine	0.55 ± 0.009	1.48 ± 0.006
Crystal-Tox	0.52 ± 0.007	1.38 ± 0.02

Chiral & Non-Chiral

Model	Macro F-Score	Weighted F-Score
Random Forest	0.19 ± 0.1	0.32 ± 0.009
K-Nearest Neighbors	0.20 ± 0.002	0.33 ± 0.002
Crystal- Olfaction	0.62 ± 0.004	0.92 ± 0.002

Model	Chiral & Non-Chiral		Enantiomer Differentiation	
	Macro F-Score	Weighted F-Score	Macro F-Score	Weighted F-Score
Random Forest	0.19 ± 0.1	0.32 ± 0.009	0.069 ± 0.002	0.31 ± 0.003
K-Nearest Neighbors	0.20 ± 0.002	0.33 ± 0.002	0.31 ± 0.0002	0.20 ± 0.001
Crystal-Olfaction	0.62 ± 0.004	0.92 ± 0.002	0.58 ± 0.003	0.93 ± 0.002

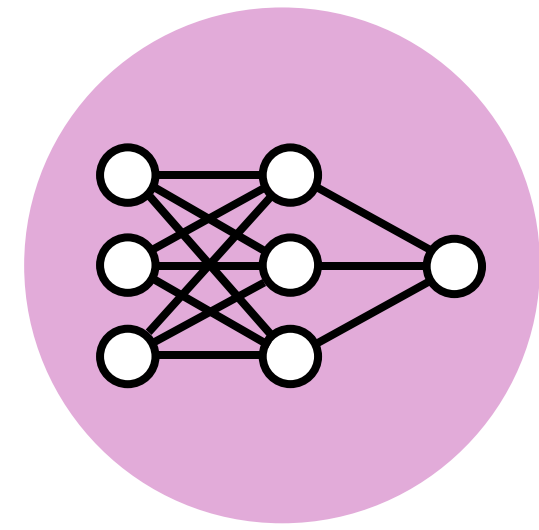




Part II:

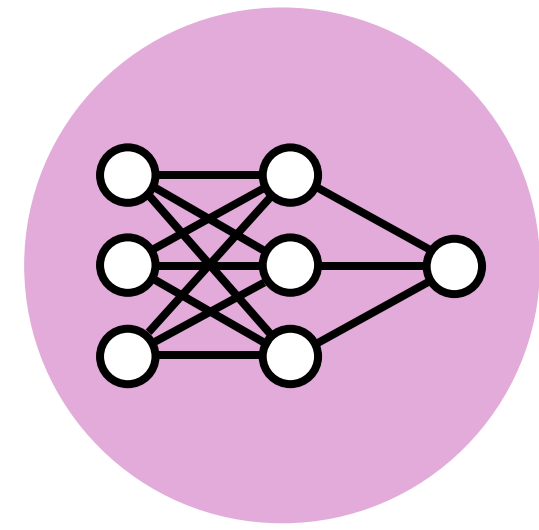
Hidden Chemical Insights from Lightweight Machine Learning

Why Lightweight ML?



Deep ML

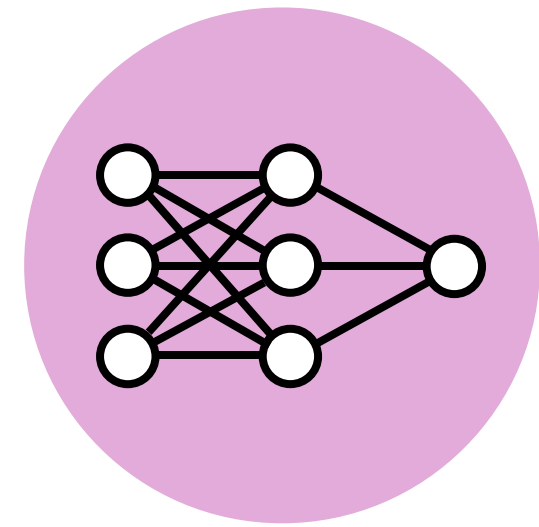
Why Lightweight ML?



Deep ML

Can get accurate predictions

Why Lightweight ML?

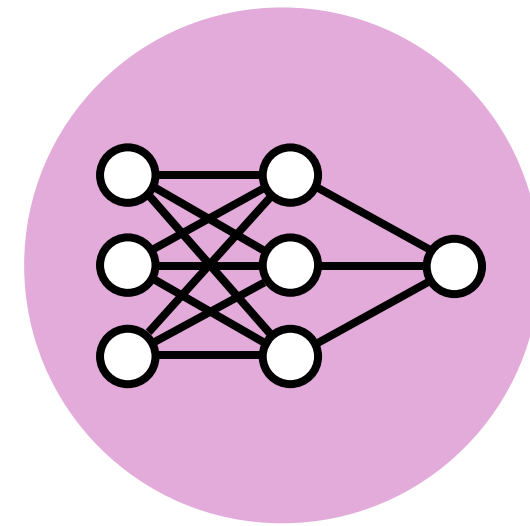


Deep ML

Can get accurate predictions

High computational resources

Why Lightweight ML?



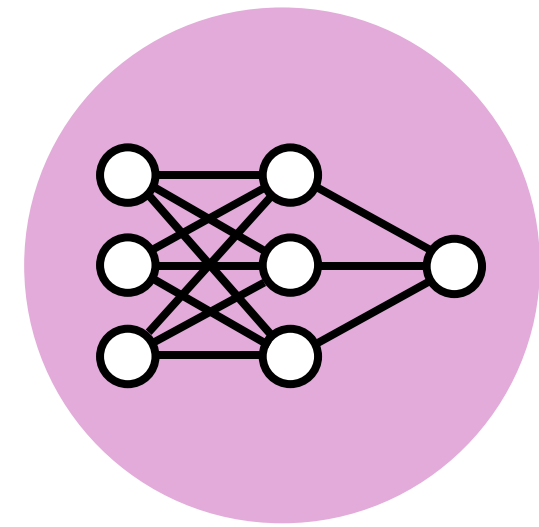
Deep ML

Can get accurate predictions

High computational resources

Careful optimization of learning architecture

Why Lightweight ML?



Deep ML

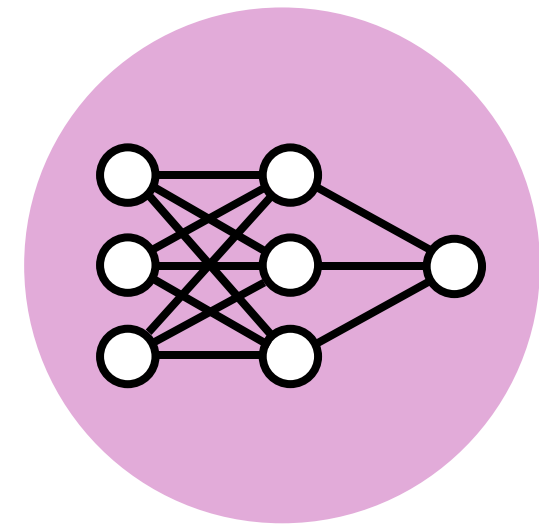
Can get accurate predictions

High computational resources

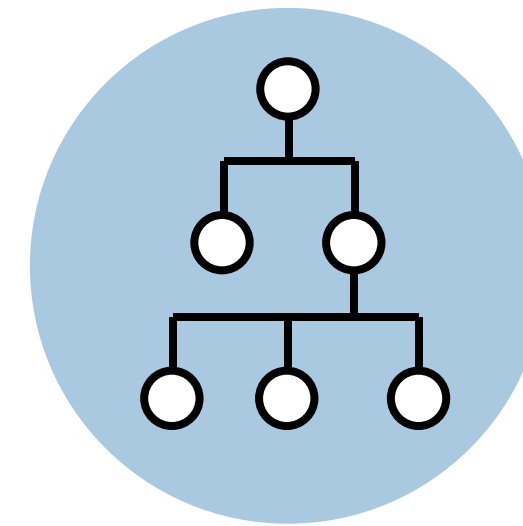
Careful optimization of learning architecture

High data requirement

Why Lightweight ML?



Deep ML



Lightweight ML

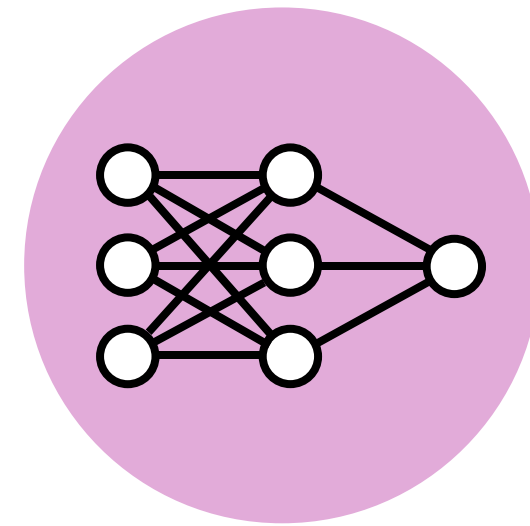
Can get accurate predictions

High computational resources

Careful optimization of learning architecture

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Why Lightweight ML?



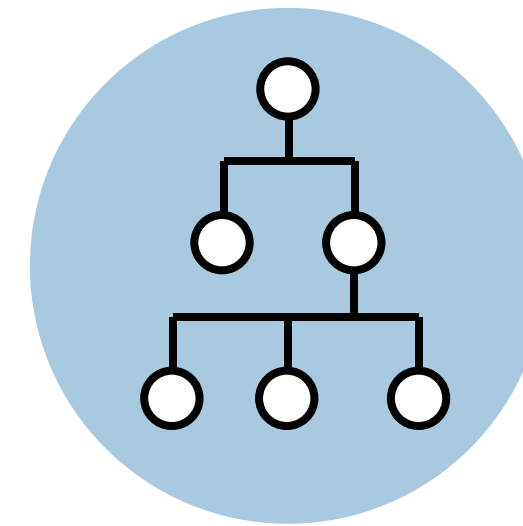
Deep ML

Can get accurate predictions

High computational resources

Careful optimization of learning architecture

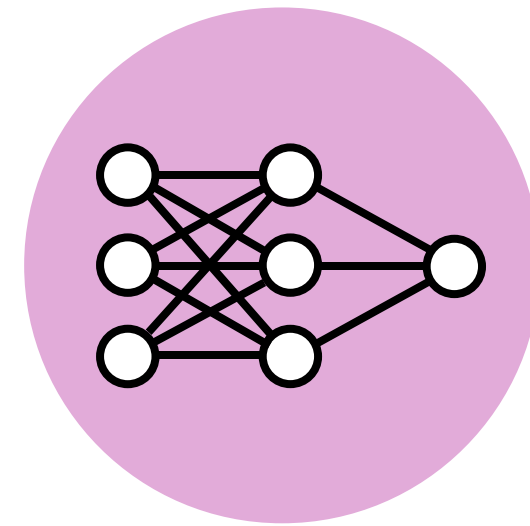
High data requirement



Lightweight ML

Run on your laptop

Why Lightweight ML?



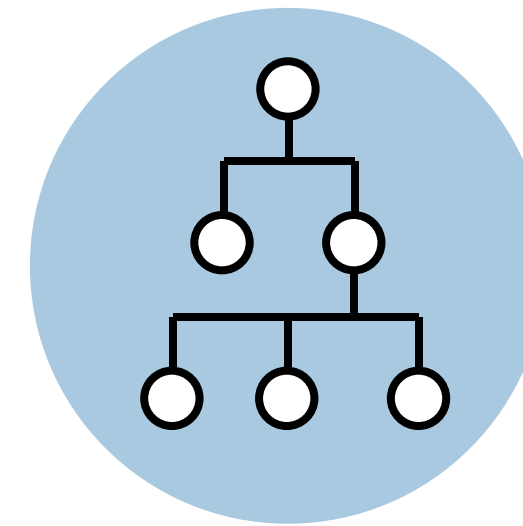
Deep ML

Can get accurate predictions

High computational resources

Careful optimization of learning architecture

High data requirement

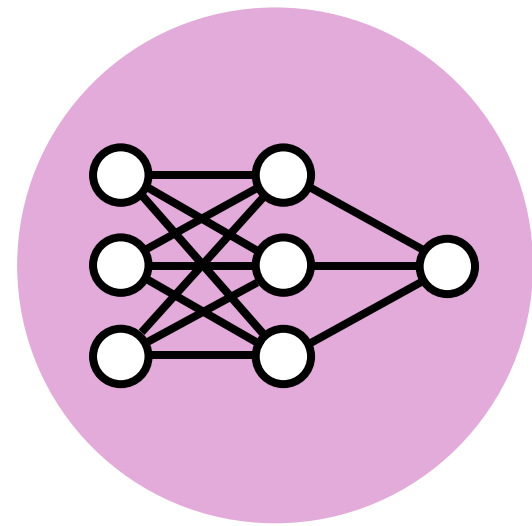


Lightweight ML

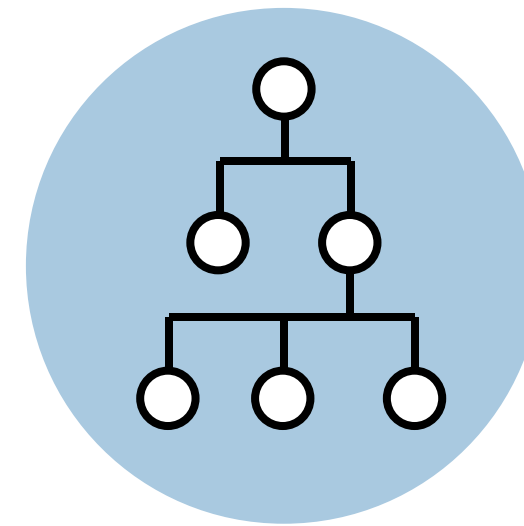
Run on your laptop

Ready to use systems

Why Lightweight ML?



Deep ML



Lightweight ML

Can get accurate predictions

High computational resources

Careful optimization of learning architecture

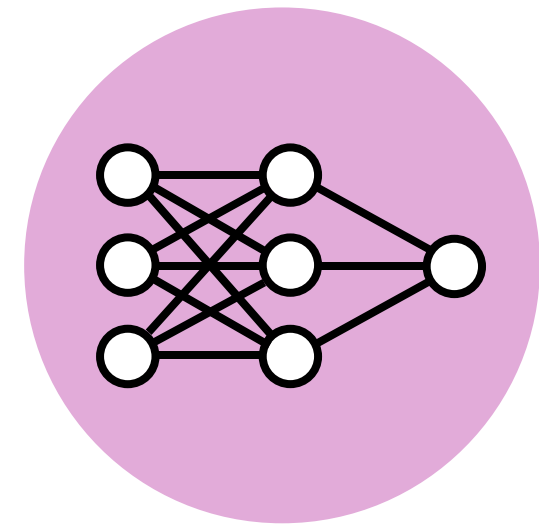
High data requirement

Run on your laptop

Ready to use systems

Lower data requirement

Why Lightweight ML?



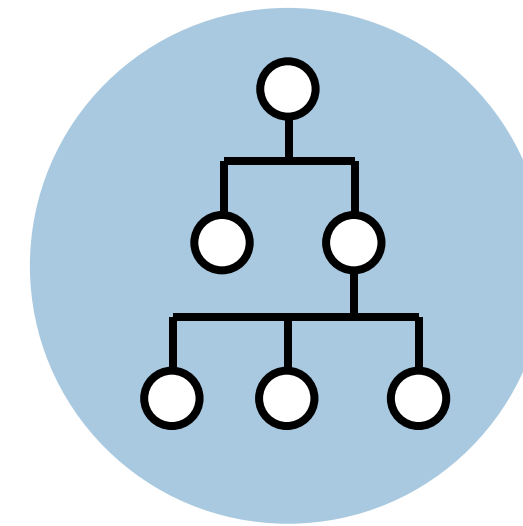
Deep ML

Can get accurate predictions

High computational resources

Careful optimization of learning architecture

High data requirement



Lightweight ML

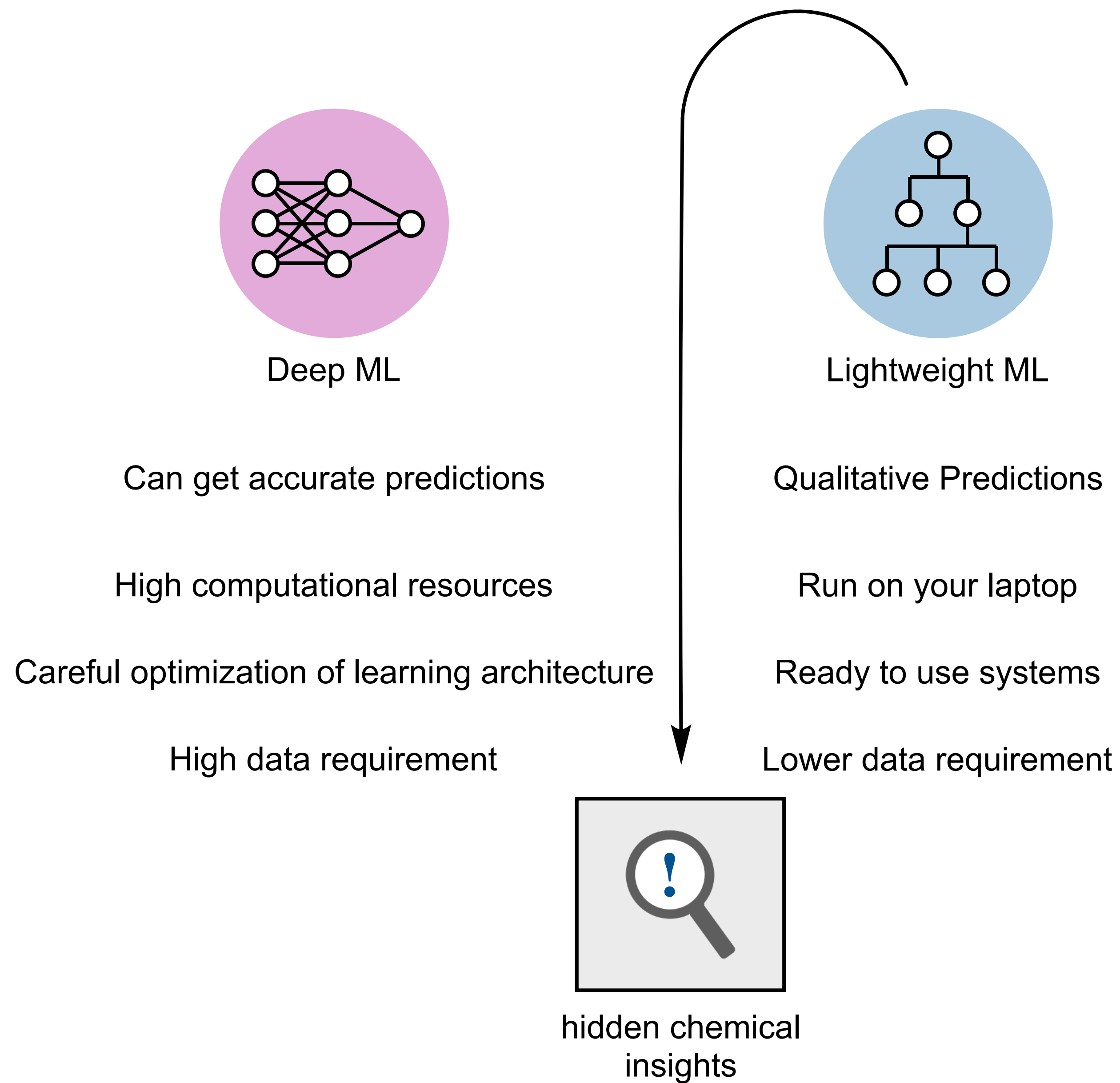
Qualitative Predictions

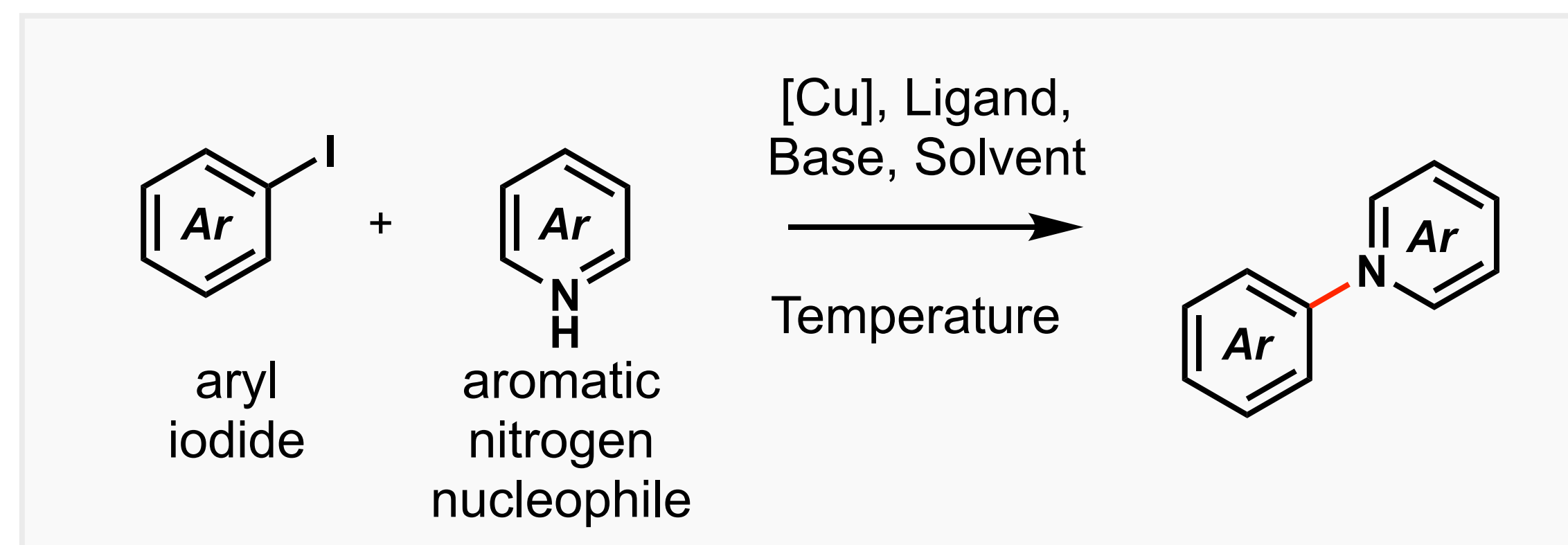
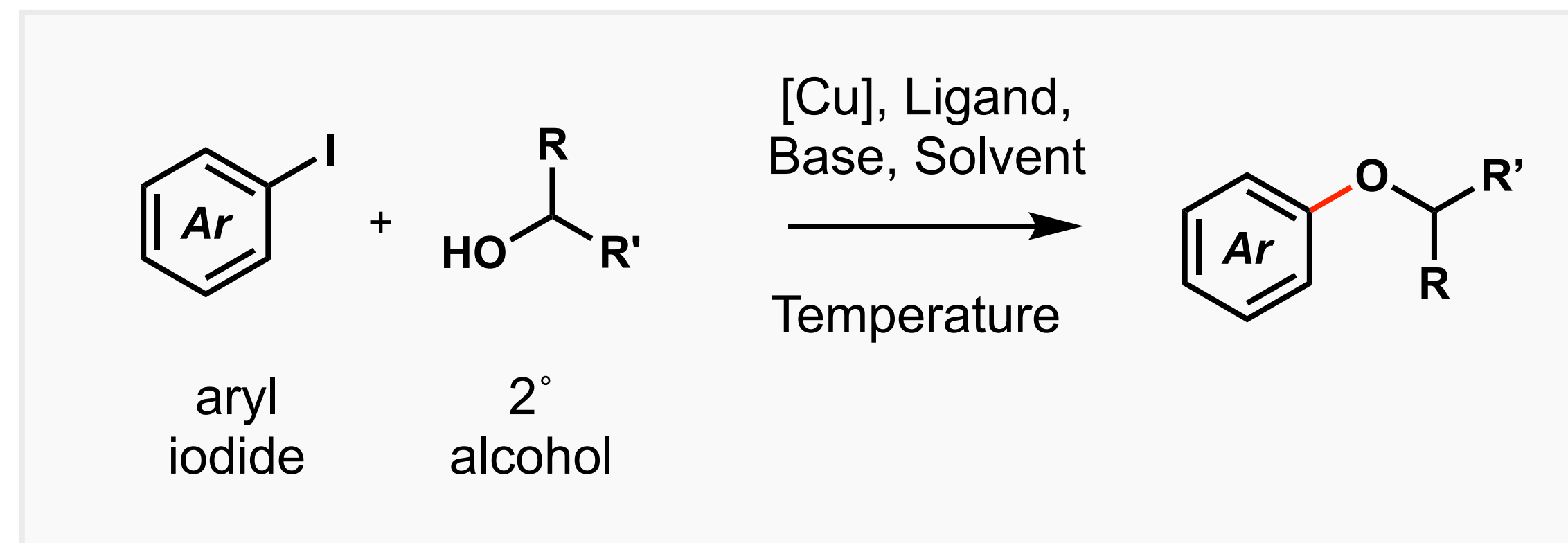
Run on your laptop

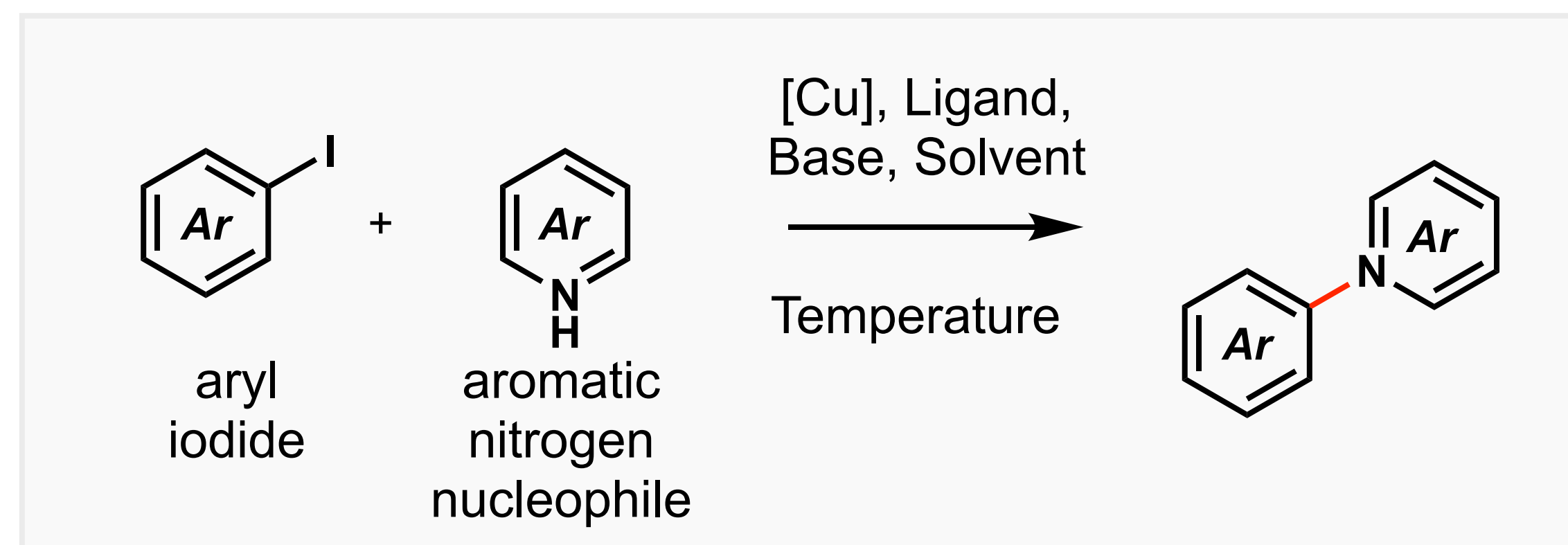
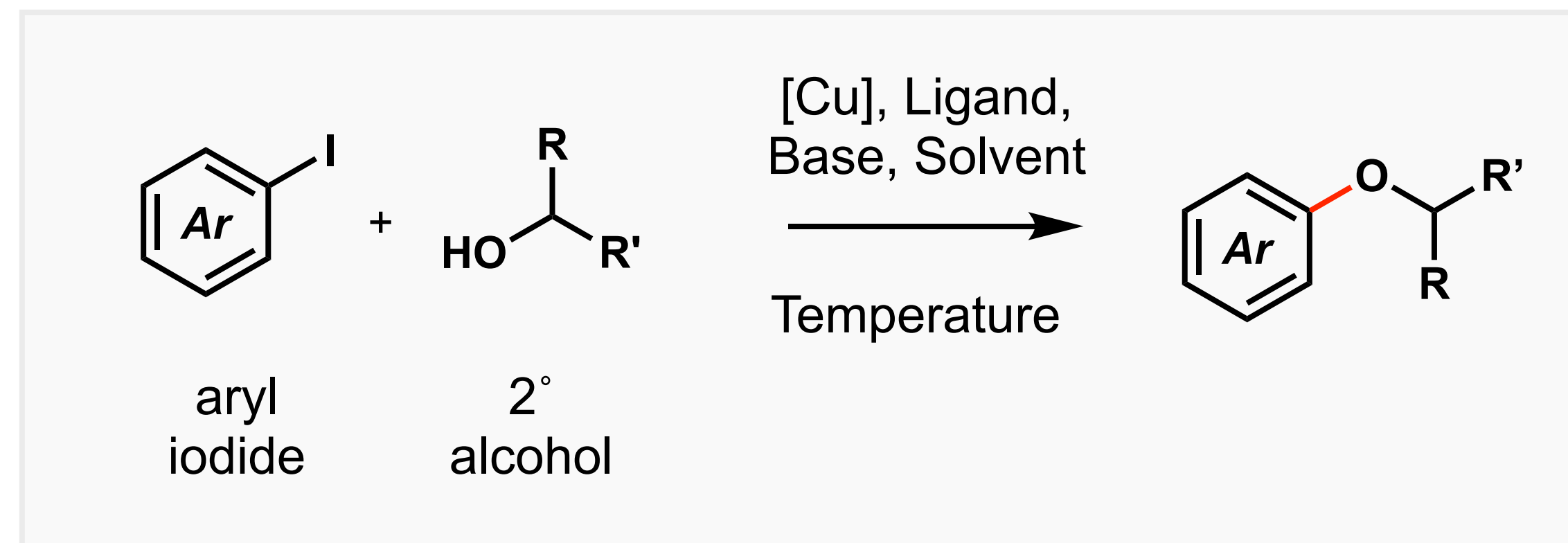
Ready to use systems

Lower data requirement

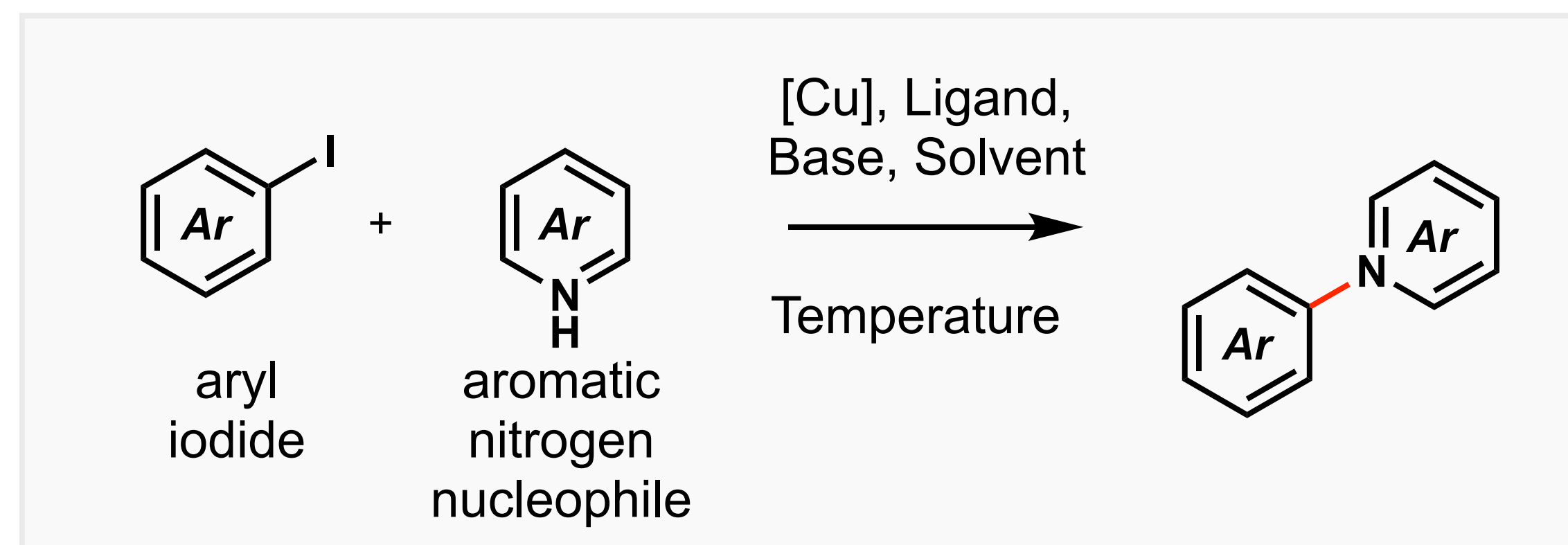
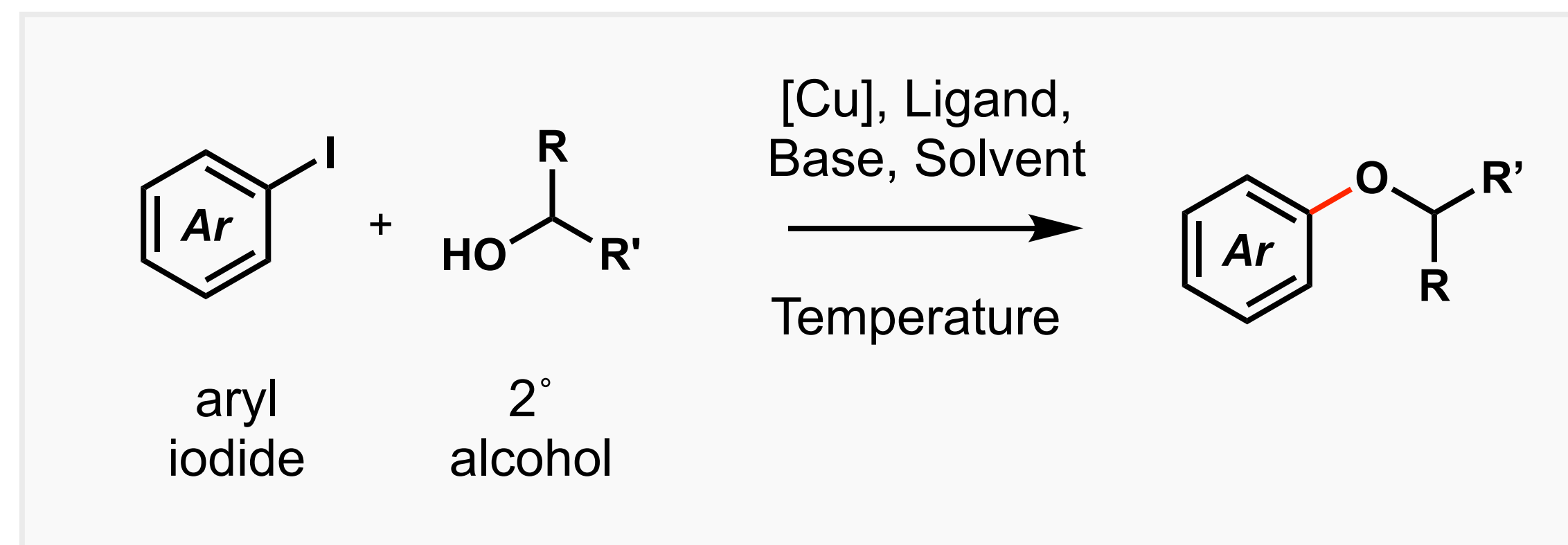
Why Lightweight ML?







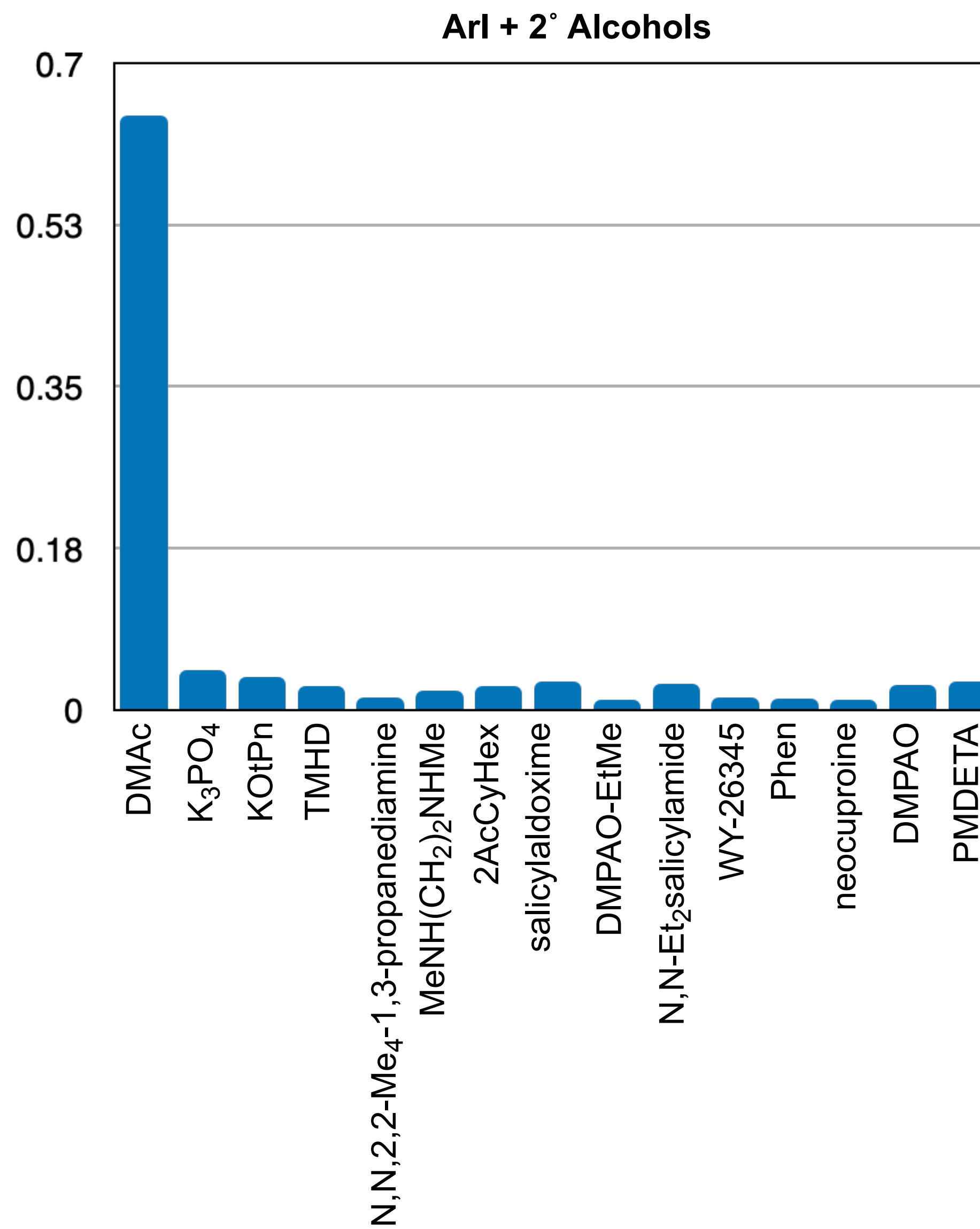
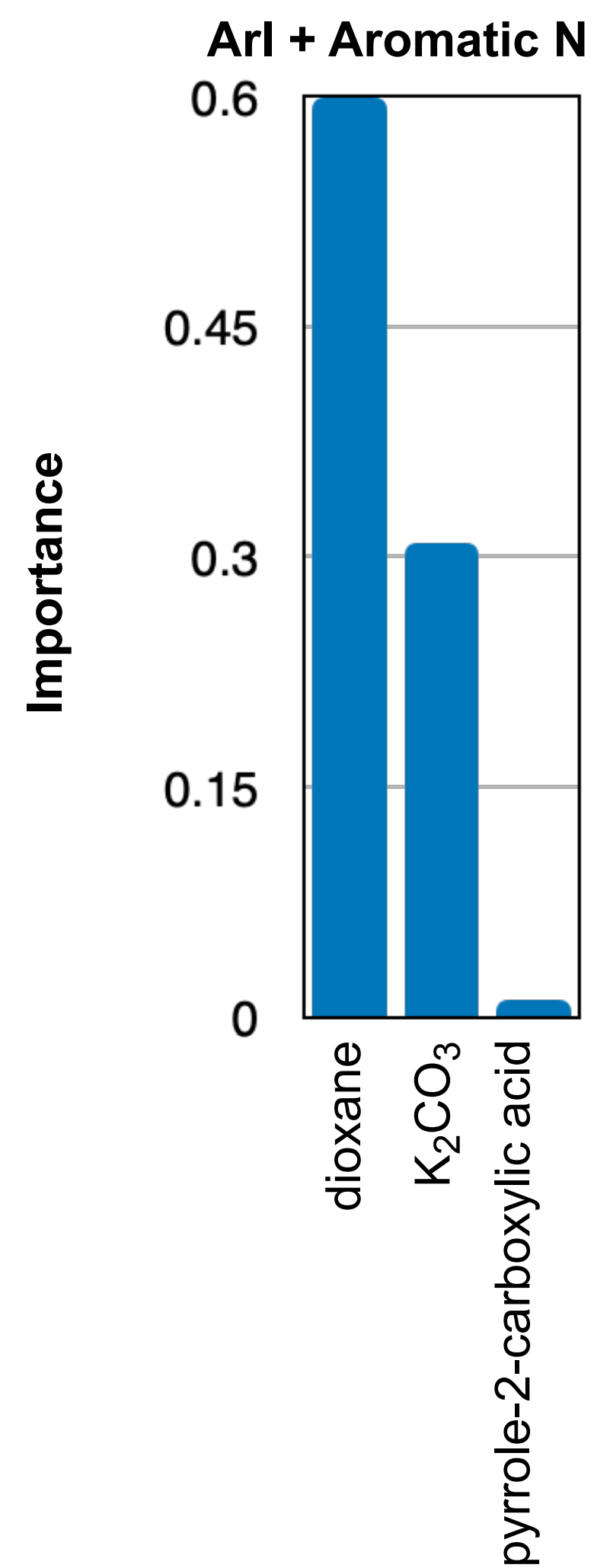
Historical data



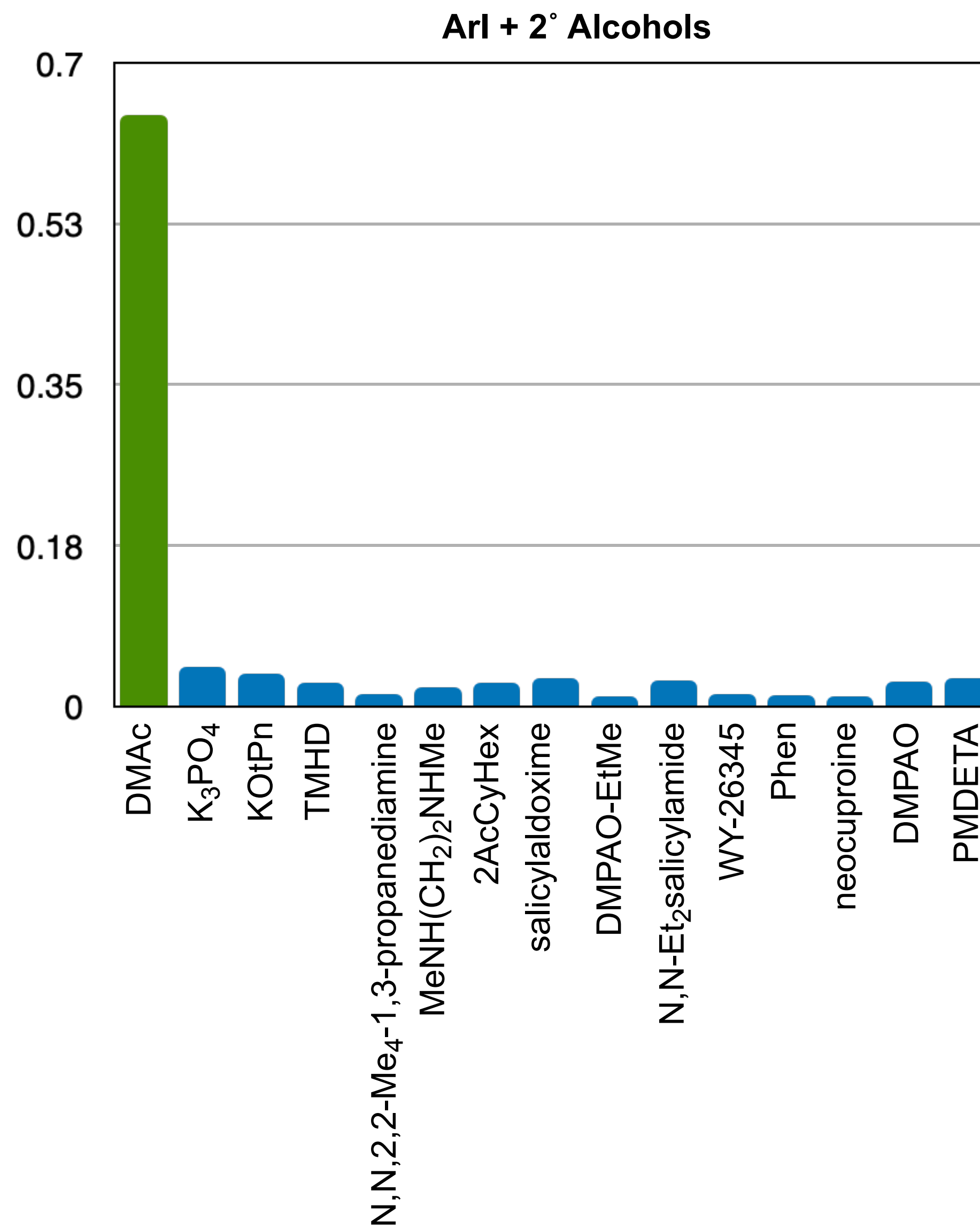
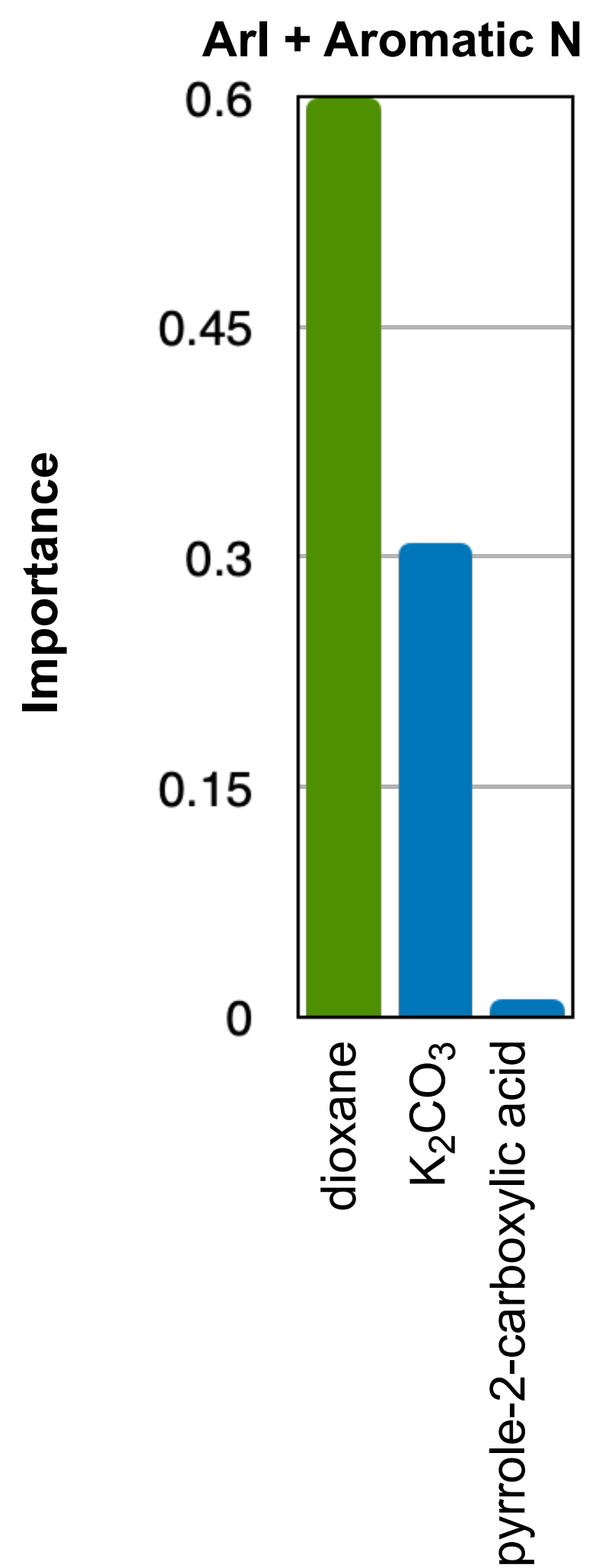
Historical data

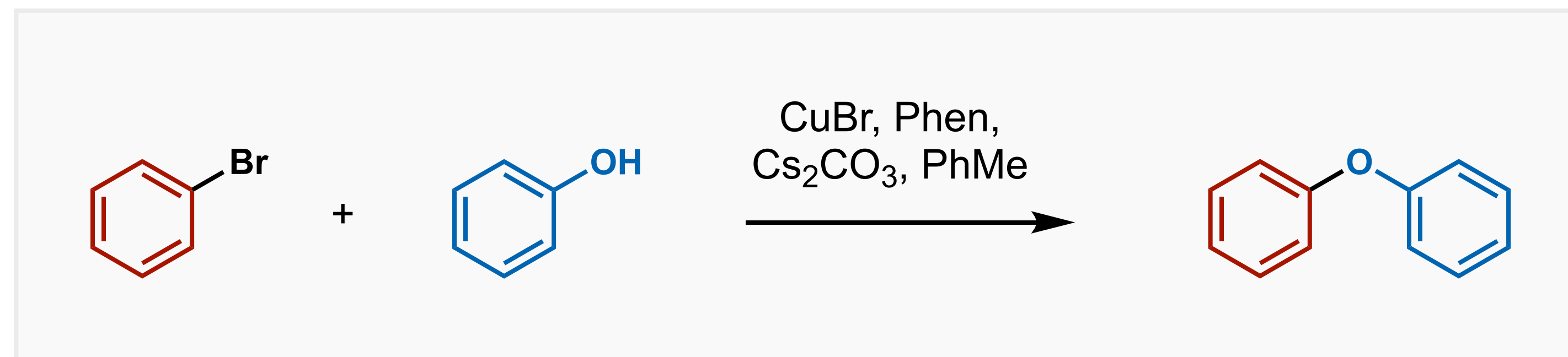
What are the important factors for reaction yield in each specific reaction class?

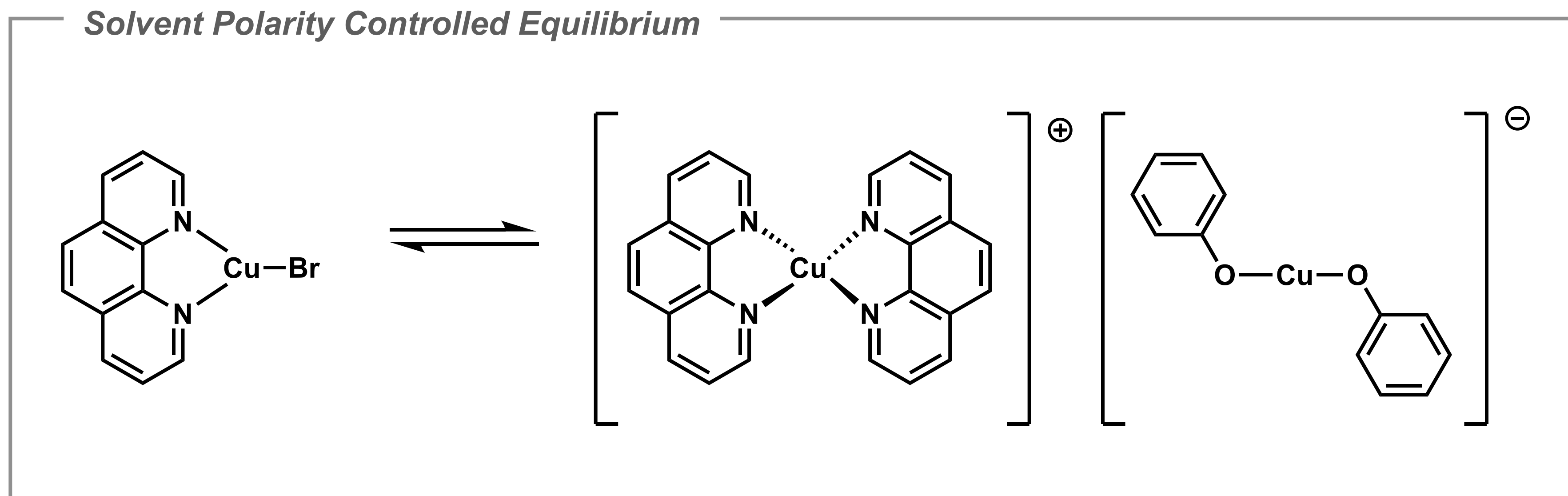
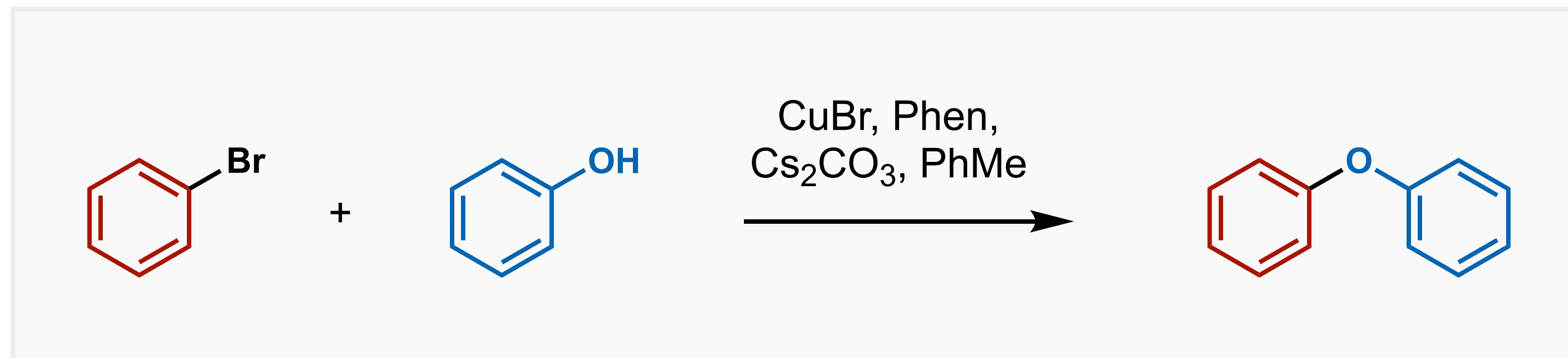
Important Features of Ullmann Condensation



Important Features of Ullmann Condensation

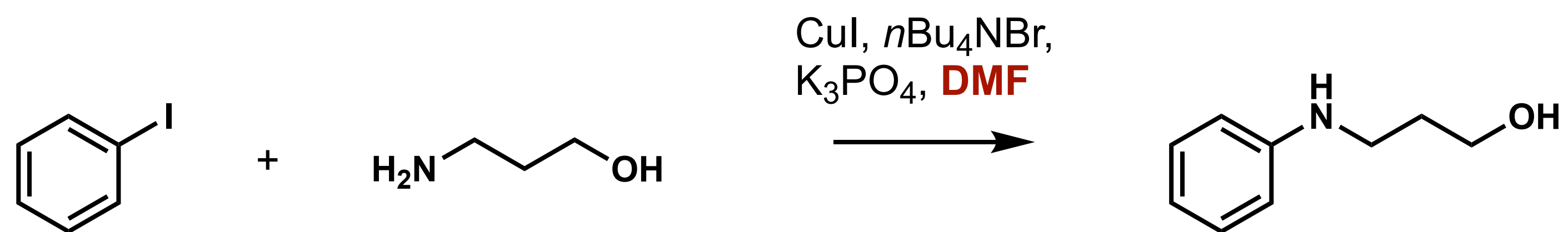




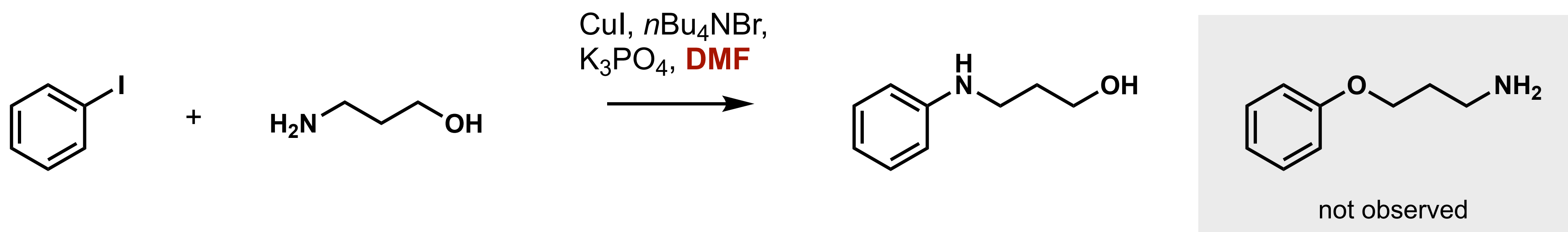


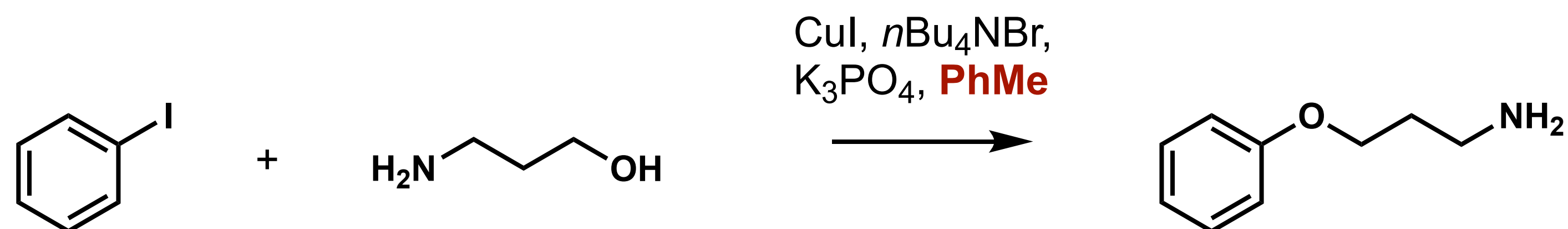
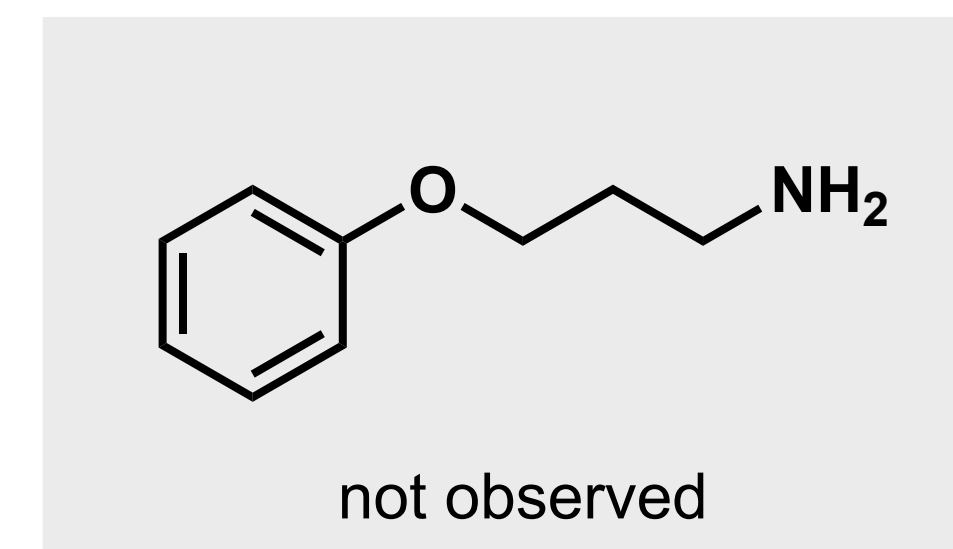
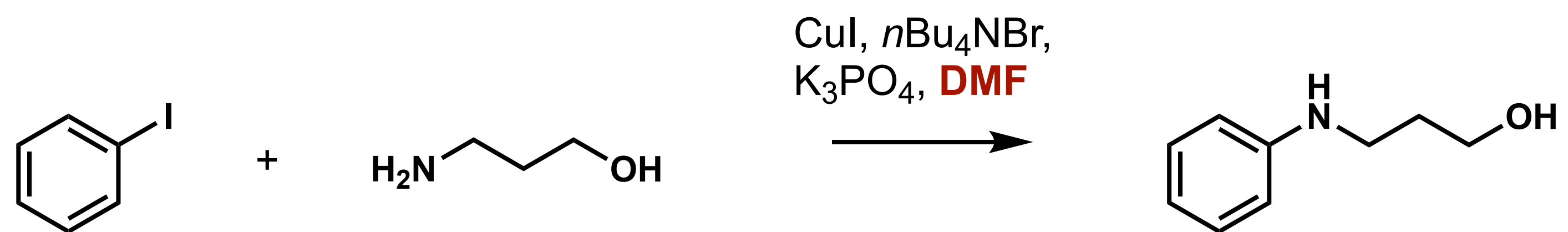
Solvent can effect active catalytic species.

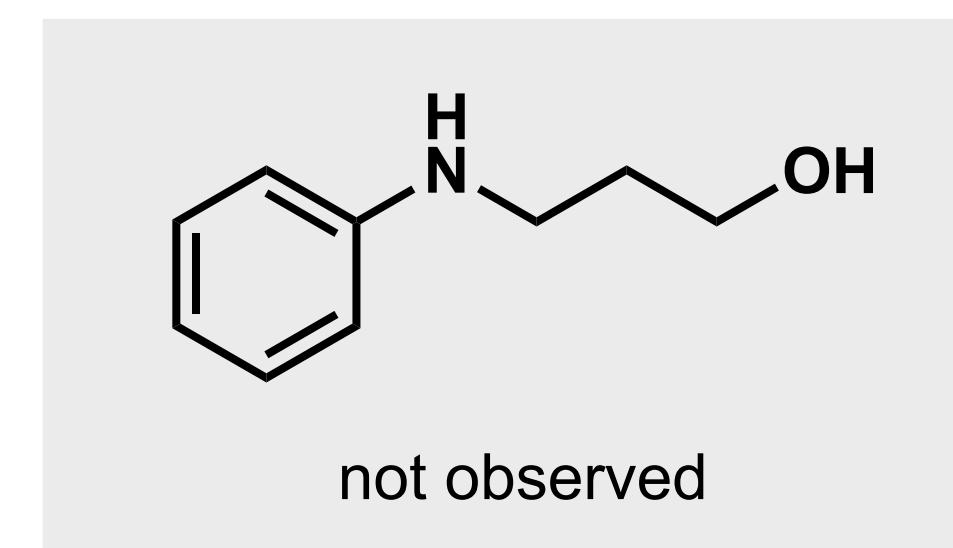
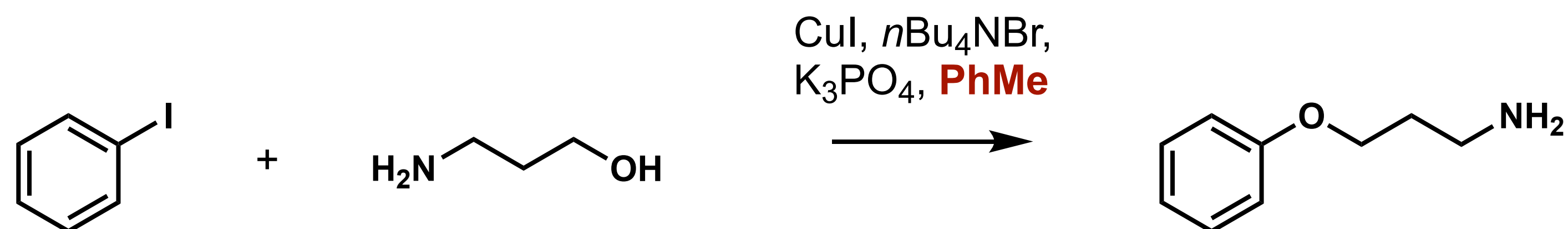
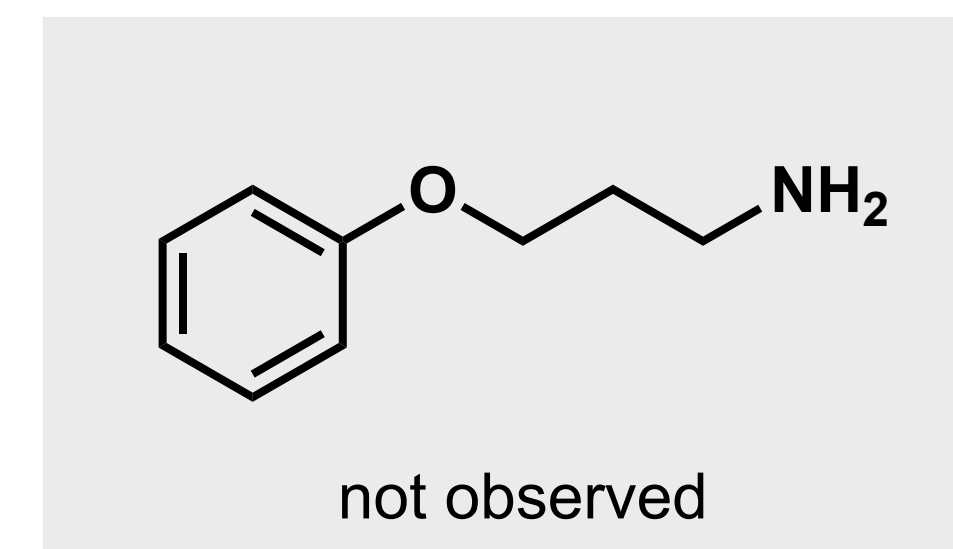
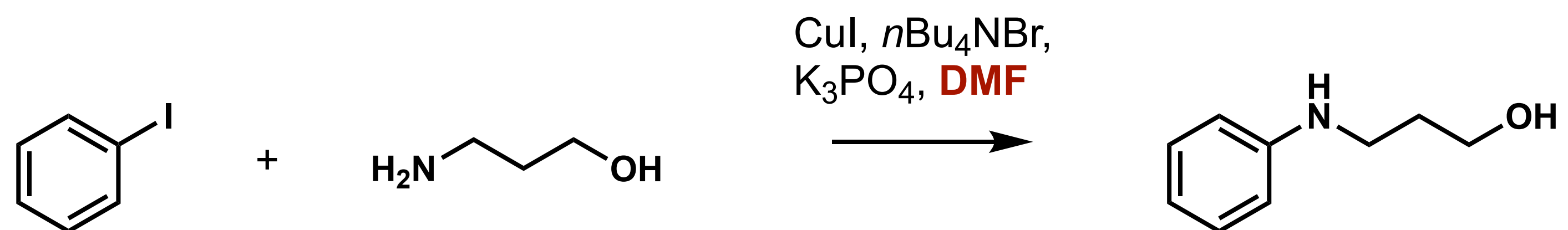
In the Literature...

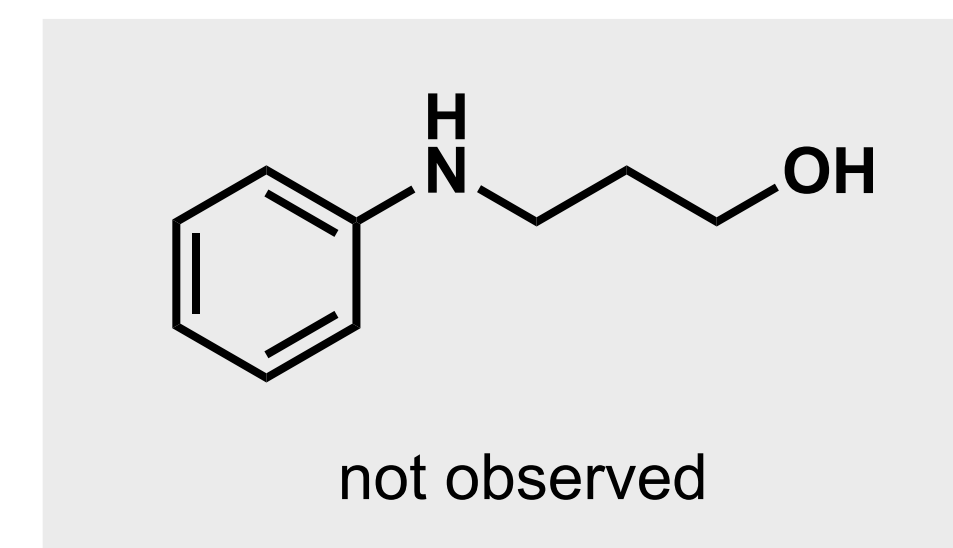
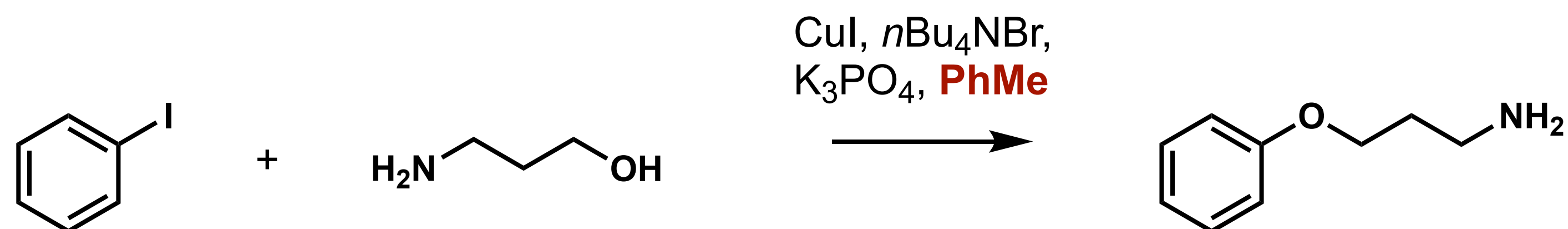
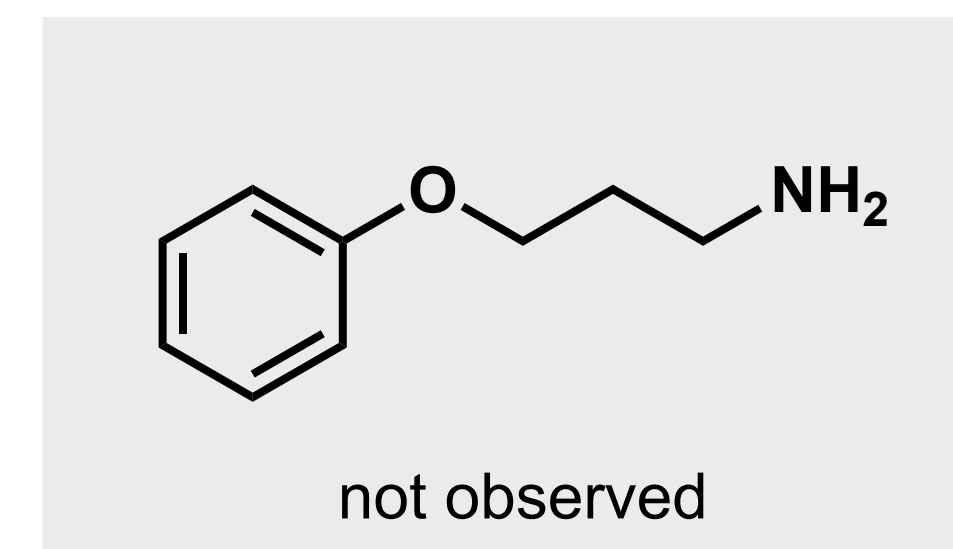
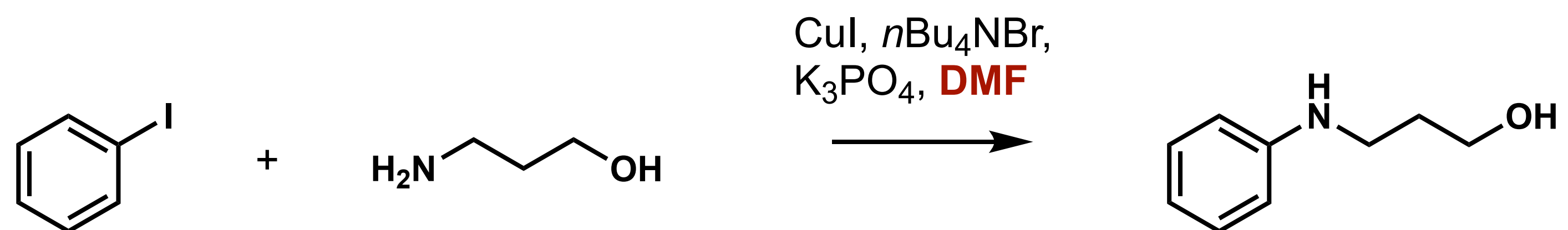


In the Literature...









No systematic review of solvent effects.

We can draw out the best solvents for a given reaction.

Acknowledgements



Lee Group

Dr. Alpha Lee

Dr. Felix Faber

Rokas Elijošius

William McCorkindale

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Dr. Simon Berritt

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Usa Reilly

Gaunt Group

Prof. Matt Gaunt

Dr Srimanta Manna

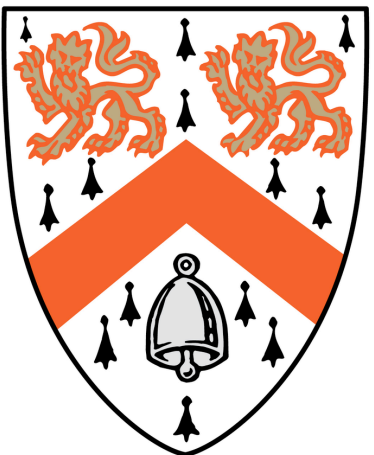
Markus Böcker

Wolfson College

Prof. Jane Clarke

Lyn Alcantara

Wolfson College Choir



Funding

Newton International Fellowship (Royal Society)

Pfizer

*Coming
Soon*

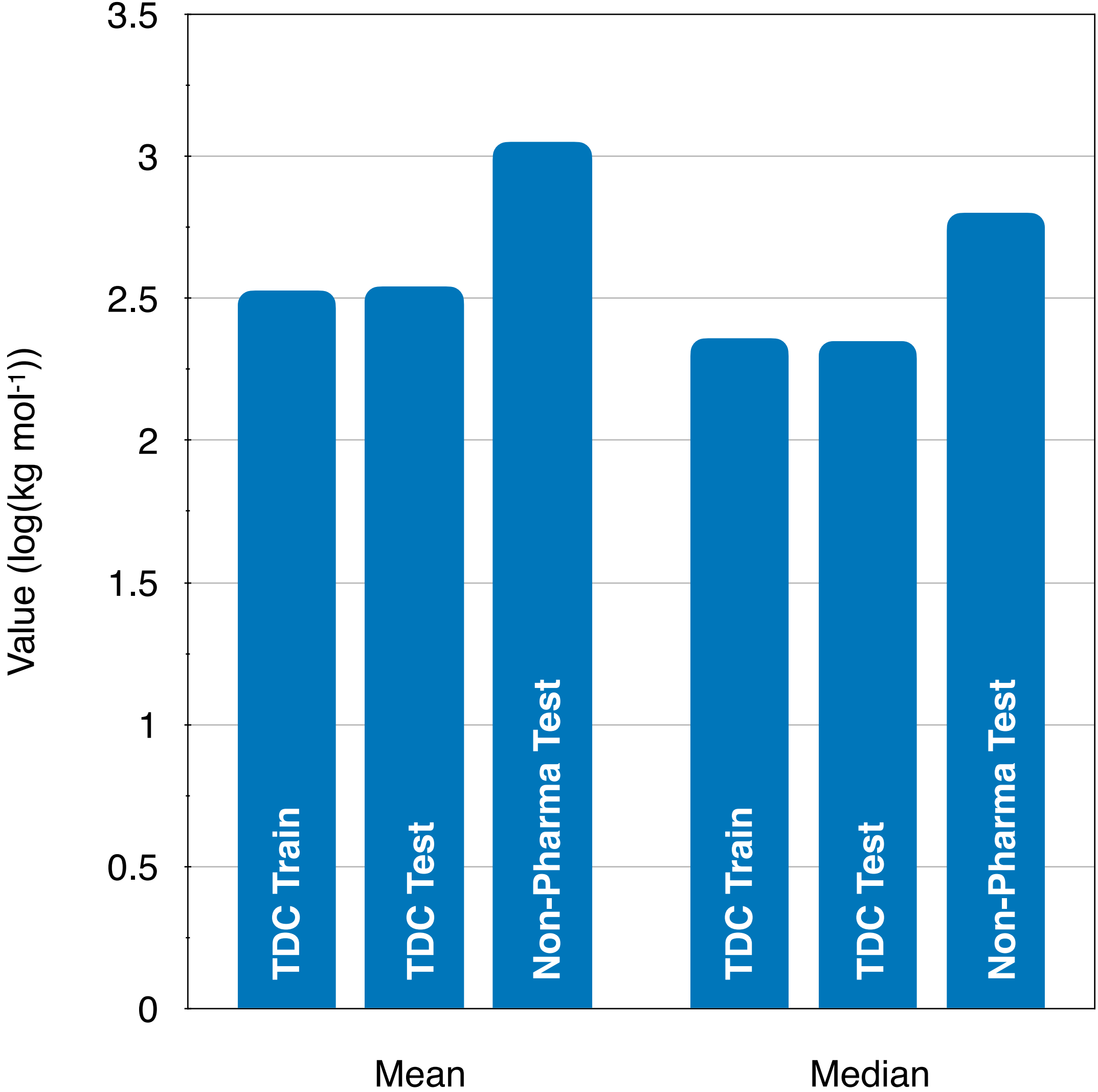


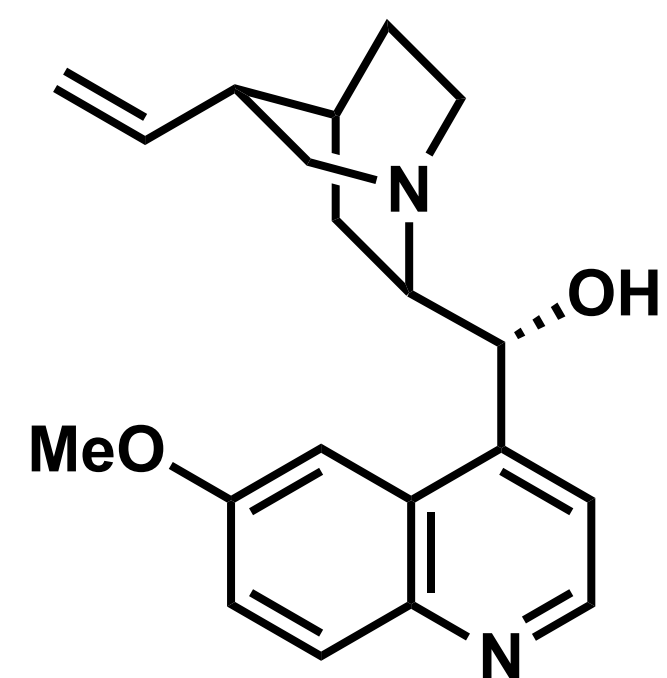
Have a Question?

esk34@cam.ac.uk



Acute Toxicity of Training and Testing Sets

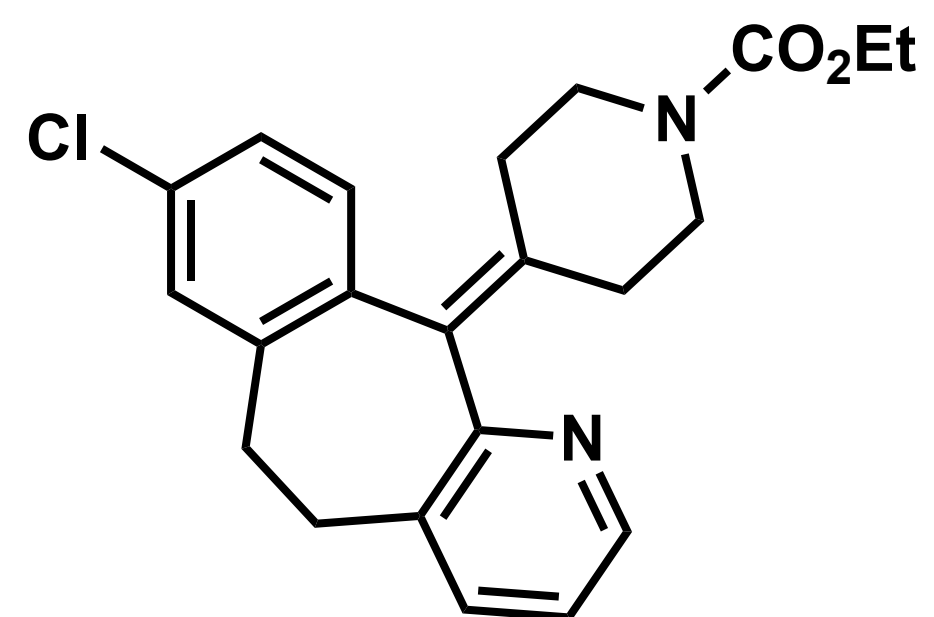




quinidine (1)

**Unique LSF
Reagents**

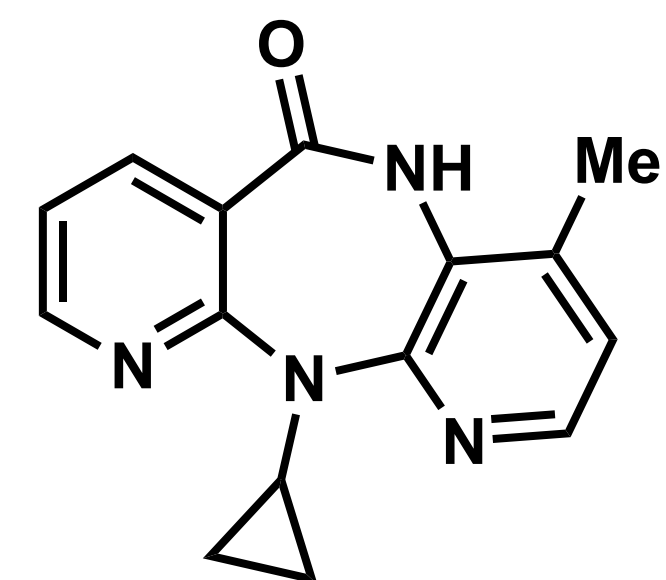
cBuBF₃K
(CF₃SO₂)₂Zn



loratadine (2)

**Unique LSF
Reagents**

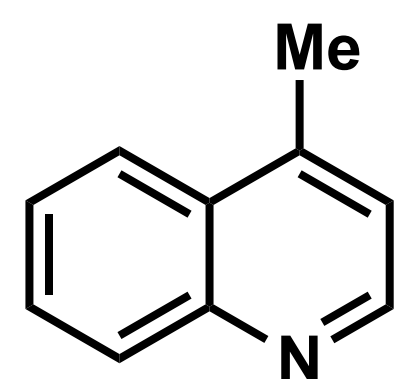
1-CF₃-cPrSO₂Na Metalloenzyme
CF₃SO₂Na (HCF₂SO₂)₂Zn
P450 HOCH₂SO₂Na



nevirapine (3)

**Unique LSF
Reagents**

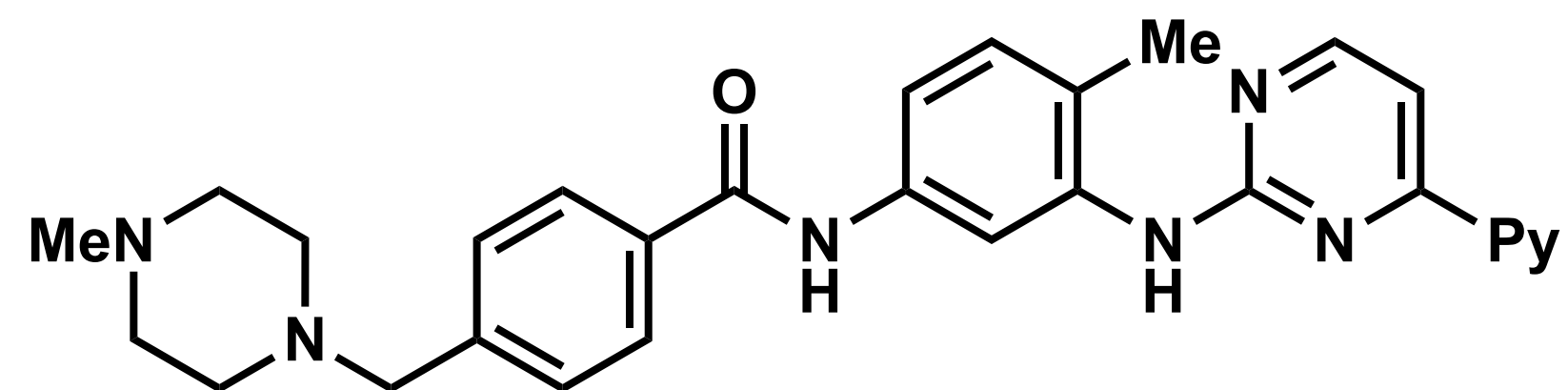
cBuBF₃K
(CF₃SO₂)₂Zn



lepidine (4)

**Unique LSF
Reagents**

NHBocCH₂BF₃K
2-Me-cPrSO₂Na
cBuBF₃K
CF₃(CH₂)₂SO₂Na

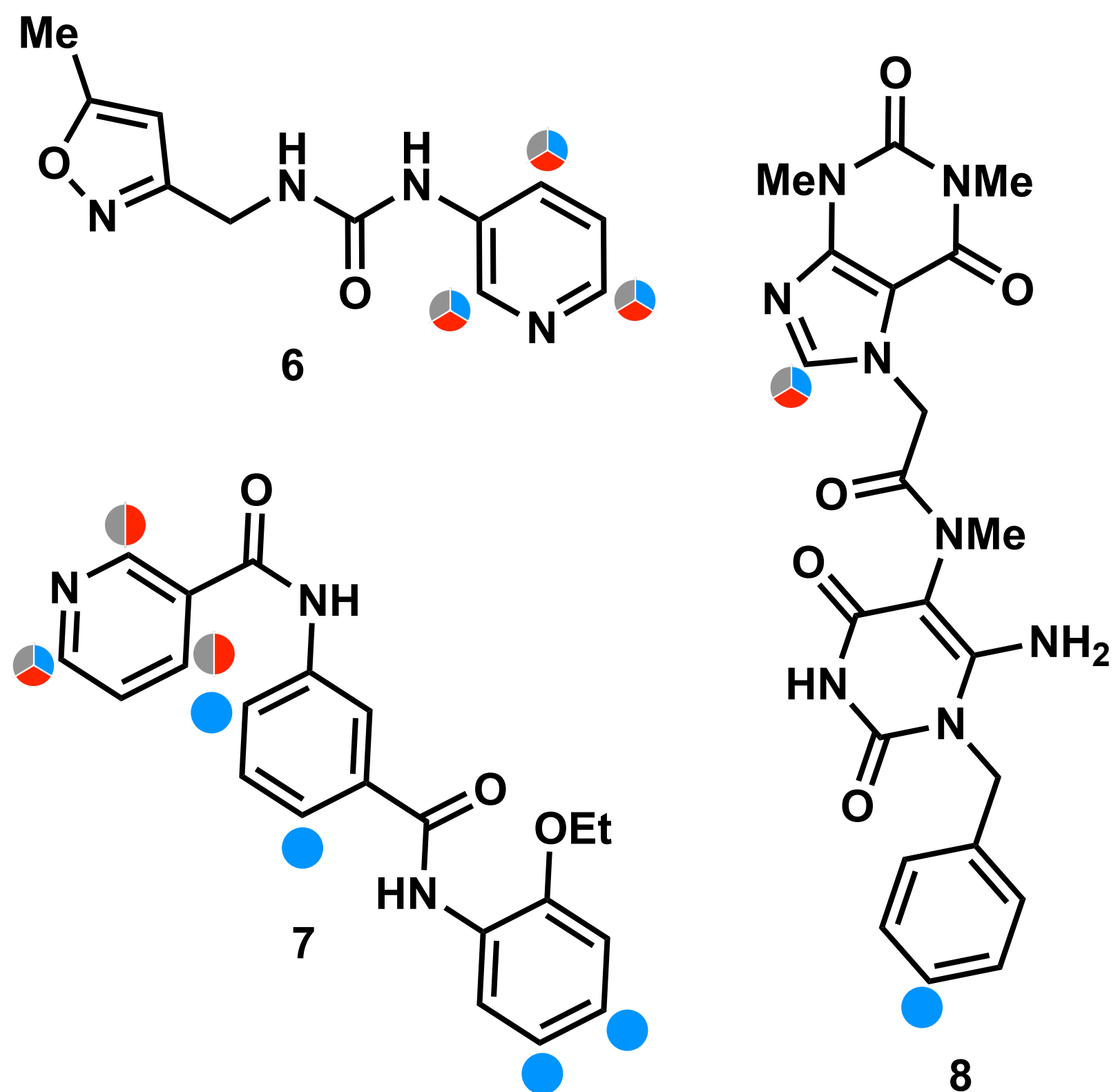


imatinib (5)

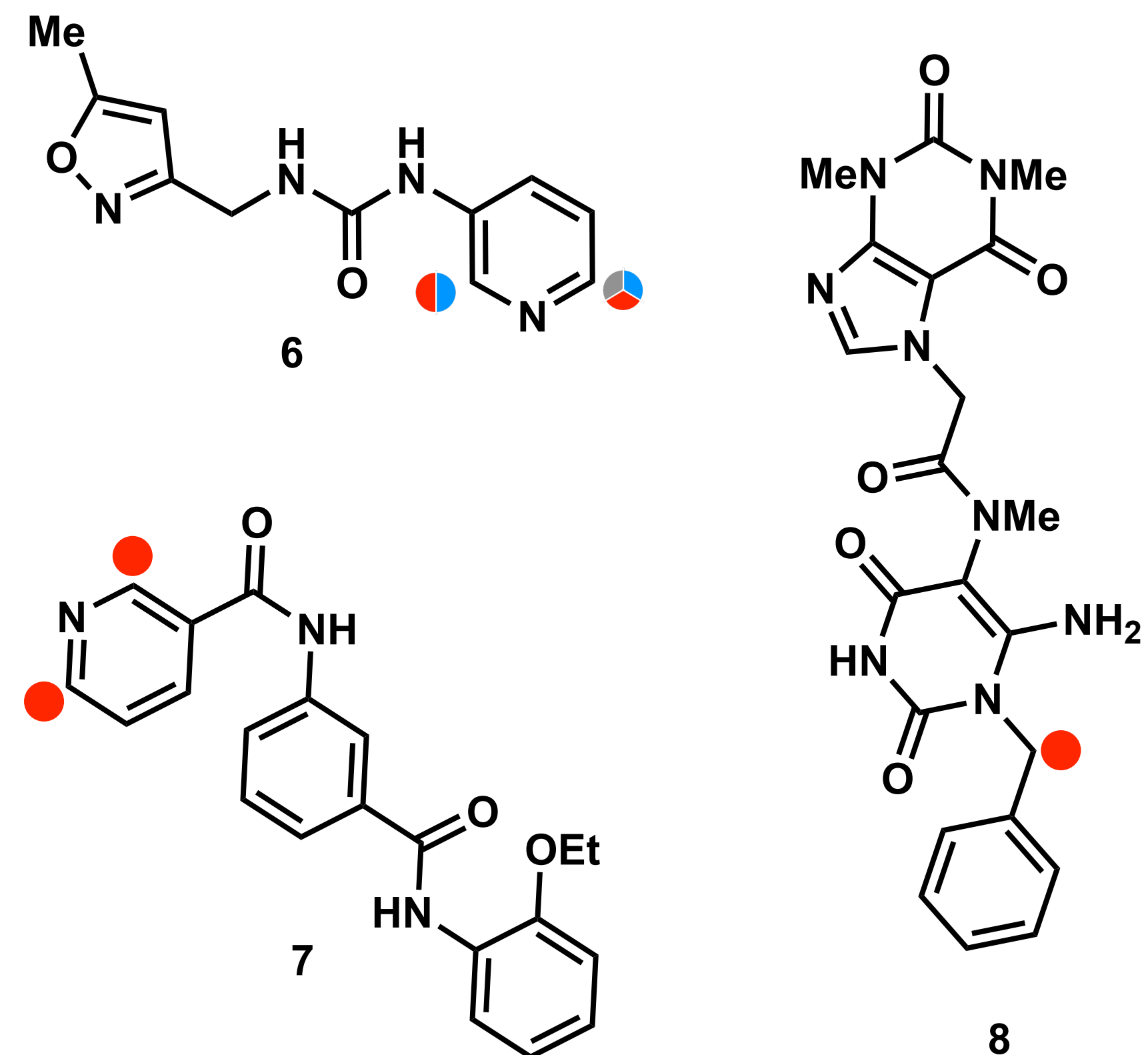
**Unique LSF
Reagents**

cBuBF₃K
HCF₂SO₂Na
(CF₃SO₂)₂Zn

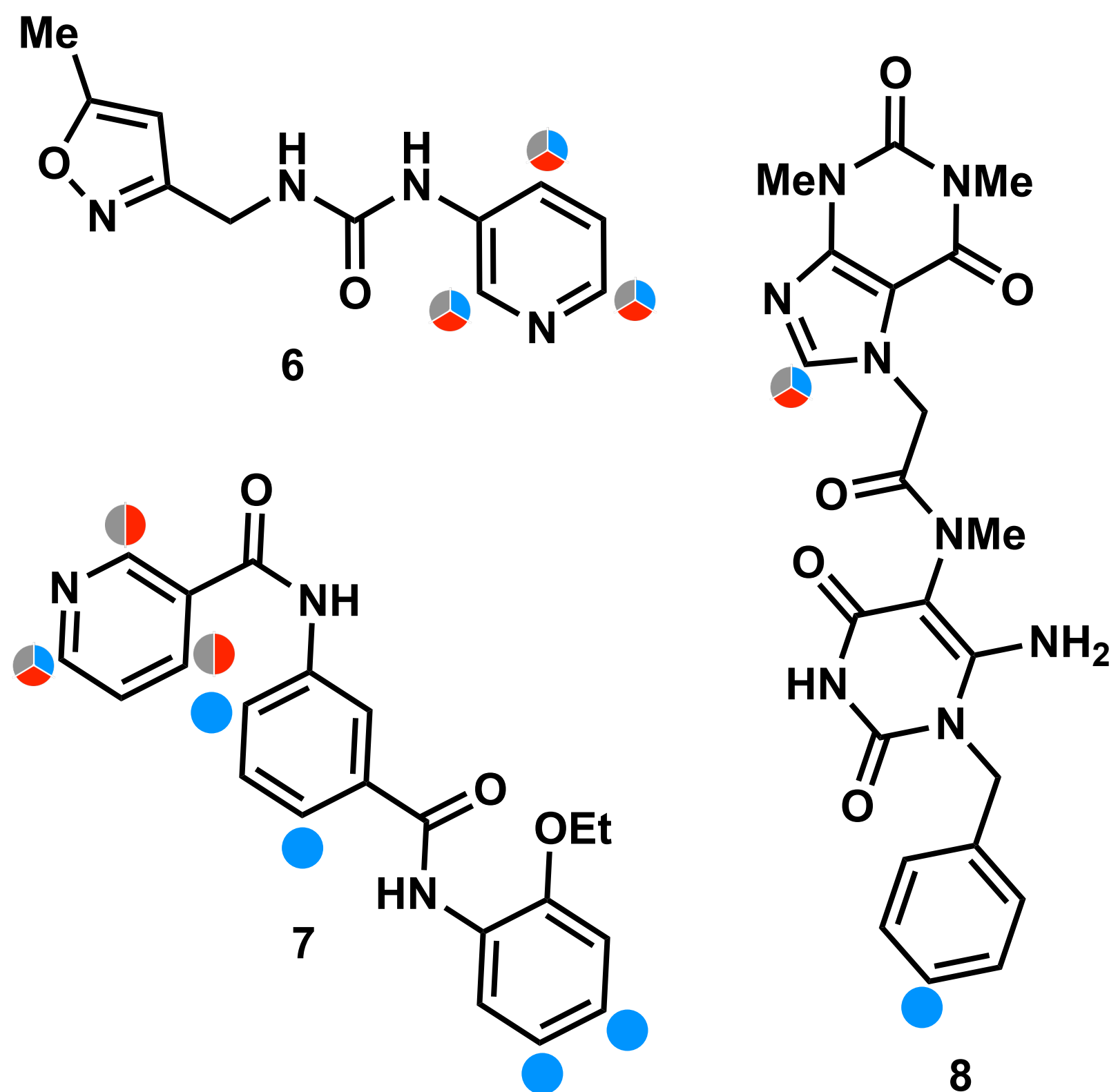
Prospective Test Set
Experimental Results



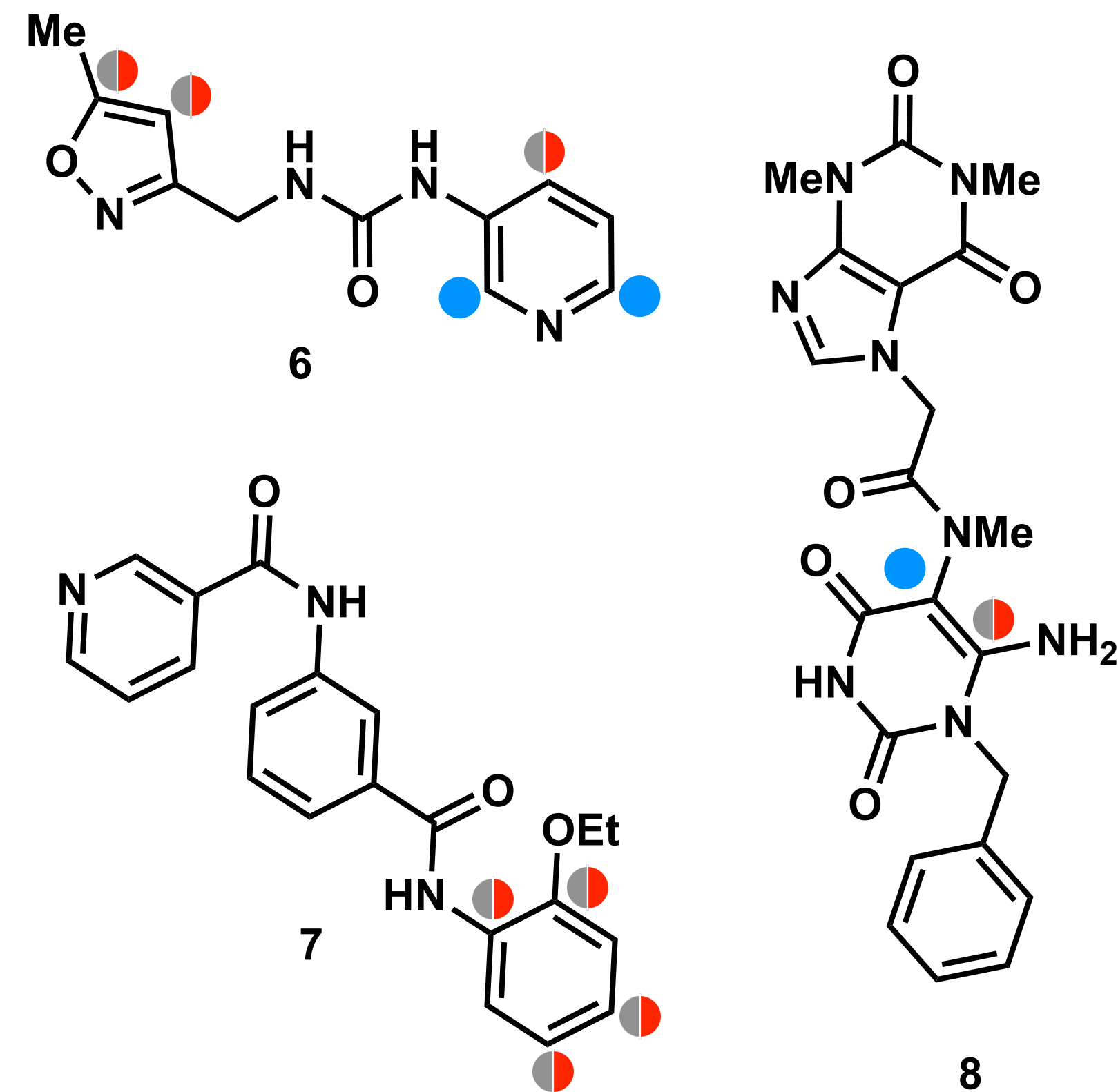
Prospective Test Set
Predictions MPNN_{LSF}



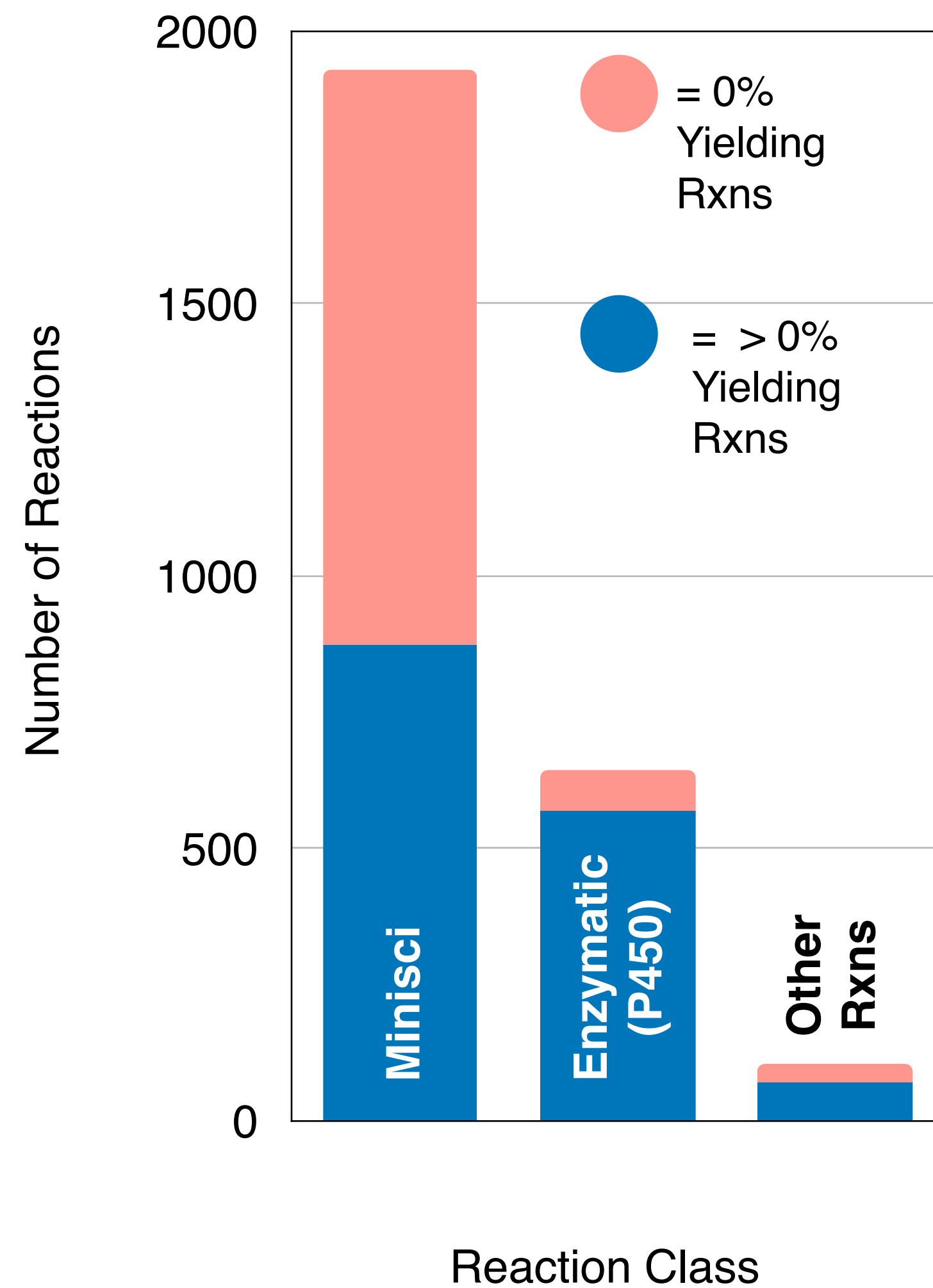
Prospective Test Set
Experimental Results



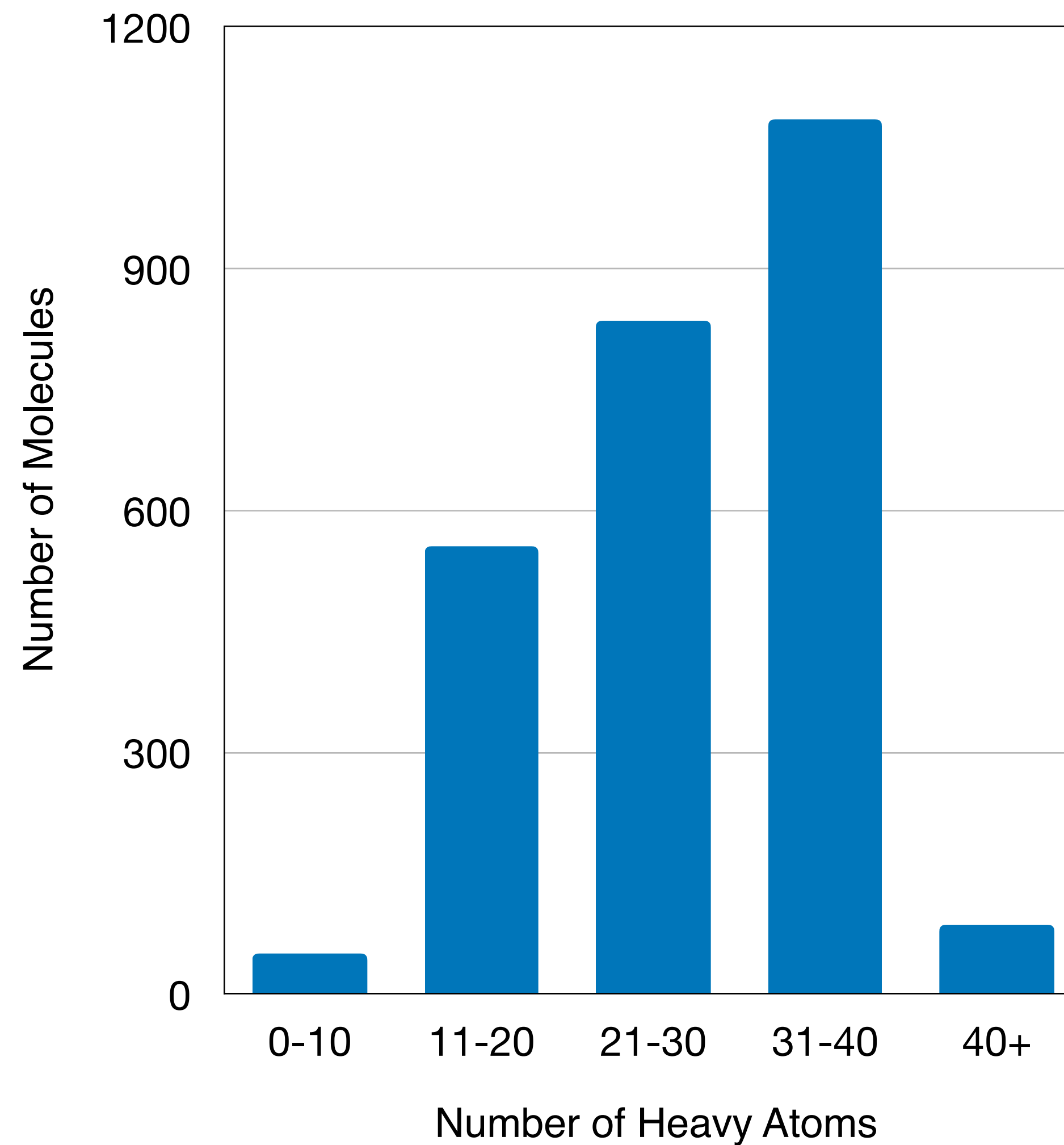
Prospective Test Set
Predictions Fukui



Dataset Breakdown: Reactions

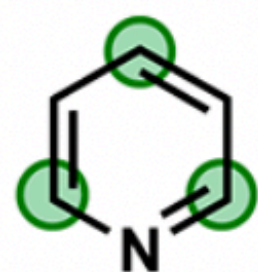


Dataset Breakdown: Molecules



STEP 1: identify innate sites

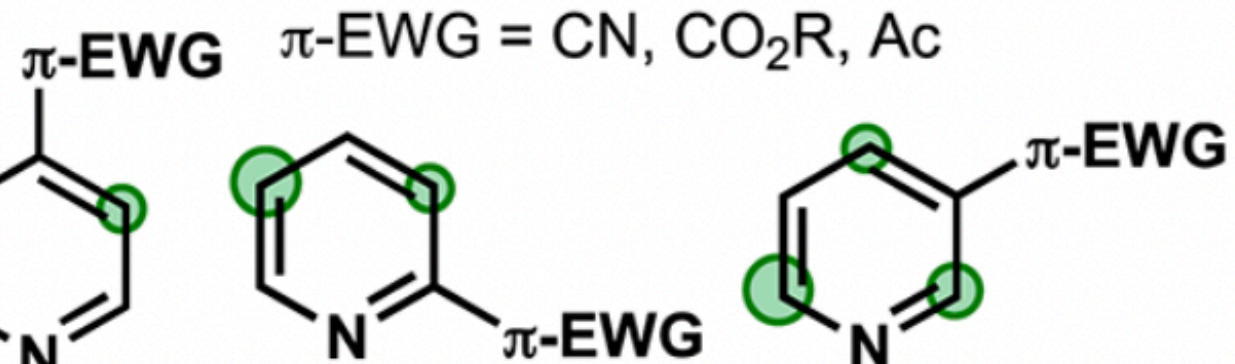
innate reactivity



activated positions:
 α and γ

STEP 2: identify conjugate sites

conjugate reactivity

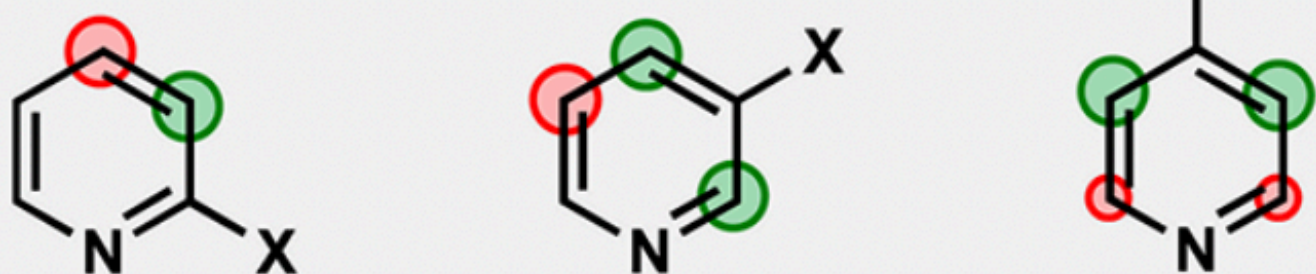


activated positions:
ortho-para to conjugating EWG

STEP 3: consider factors that modify reactivity of *activated* sites

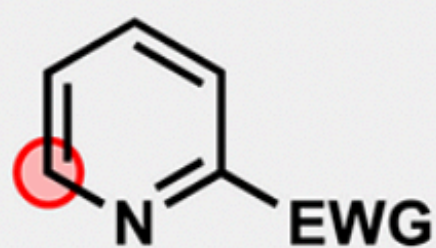
halide, alkoxy

activate *ortho*, deactivate *meta*



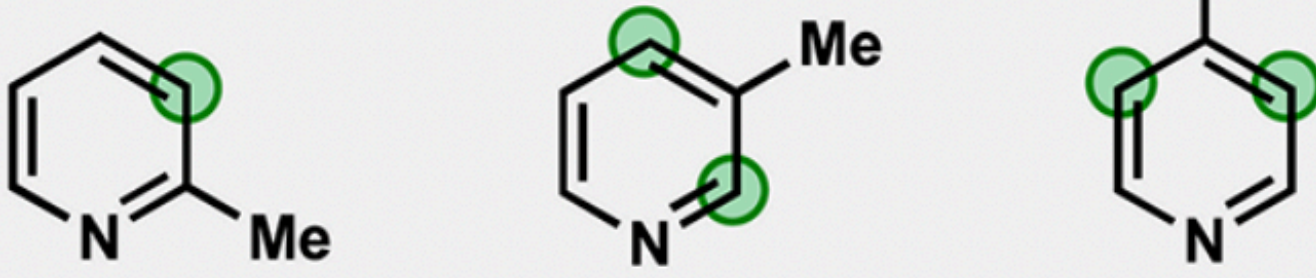
C2 EWG

deactivate C6



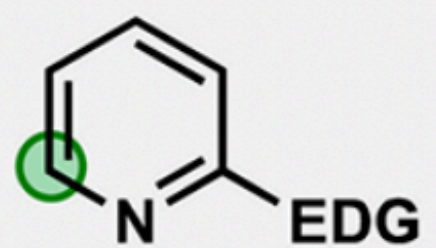
methyl

activate *ortho*







C2 EDG

activate C6




STEP 4: add effects and decide on conditions

 promote innate reactivity
 reduce conjugate reactivity


 reduce innate reactivity
 promote conjugate reactivity

solvent + acid

 increase reactivity for
electron-rich systems

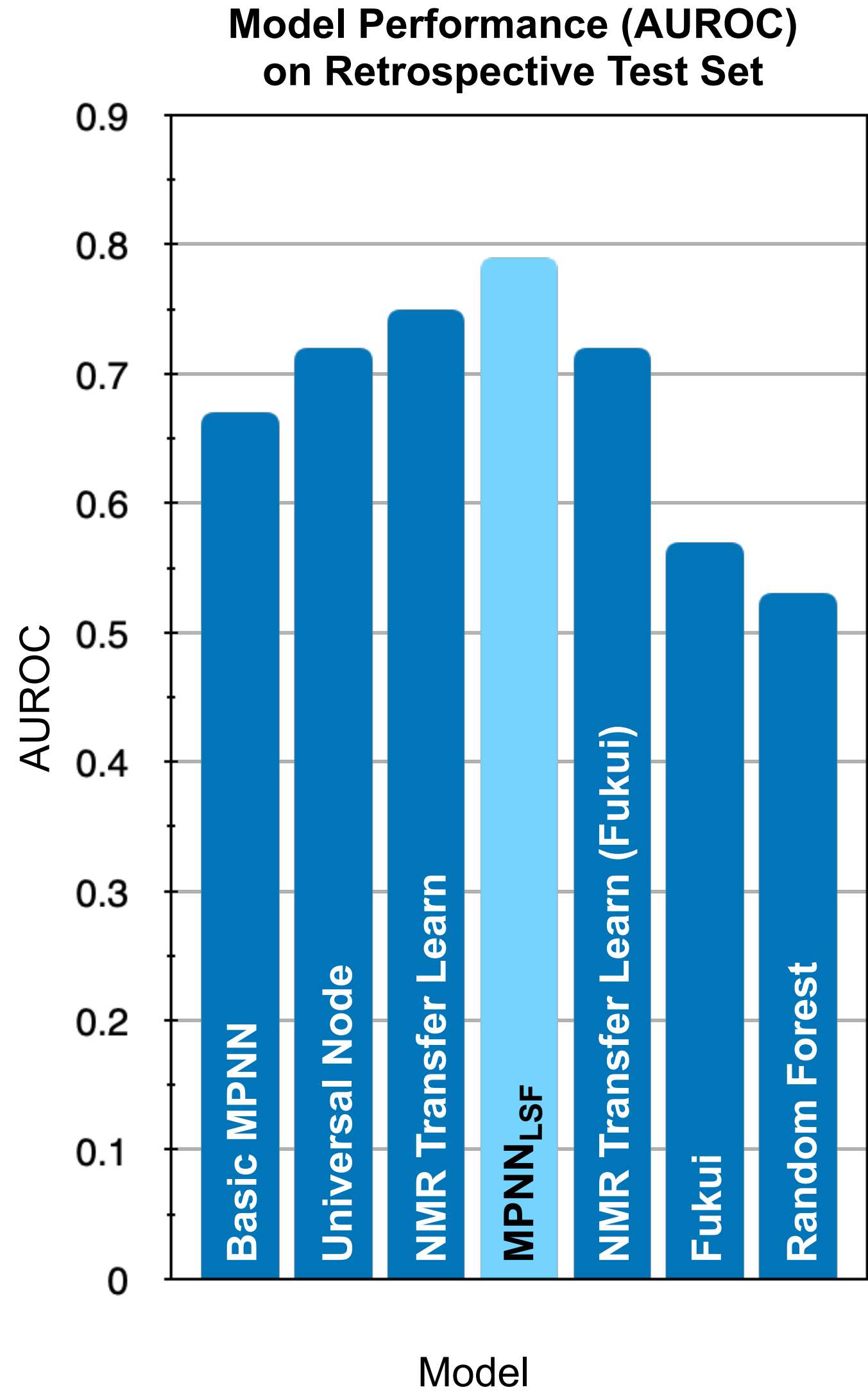
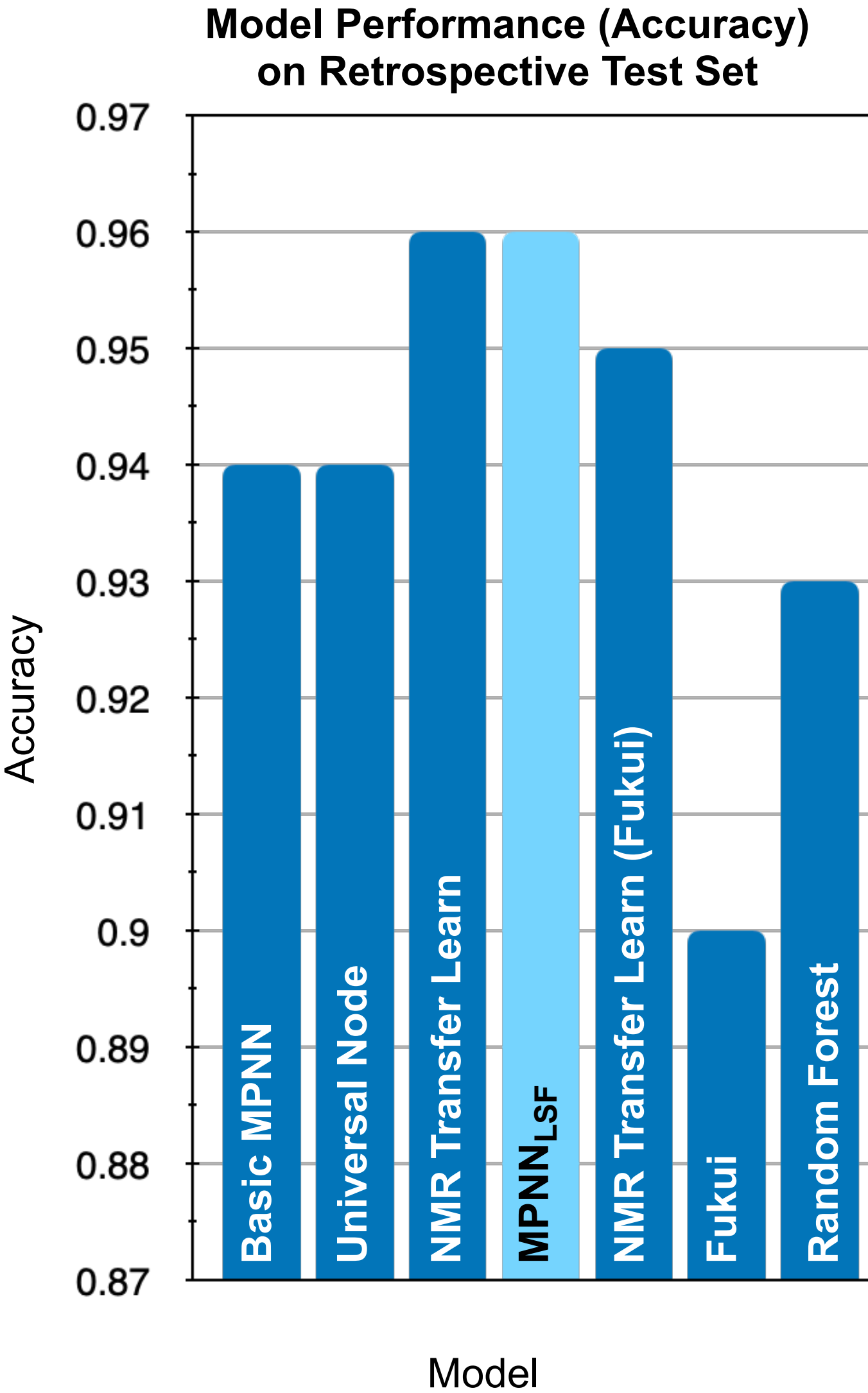
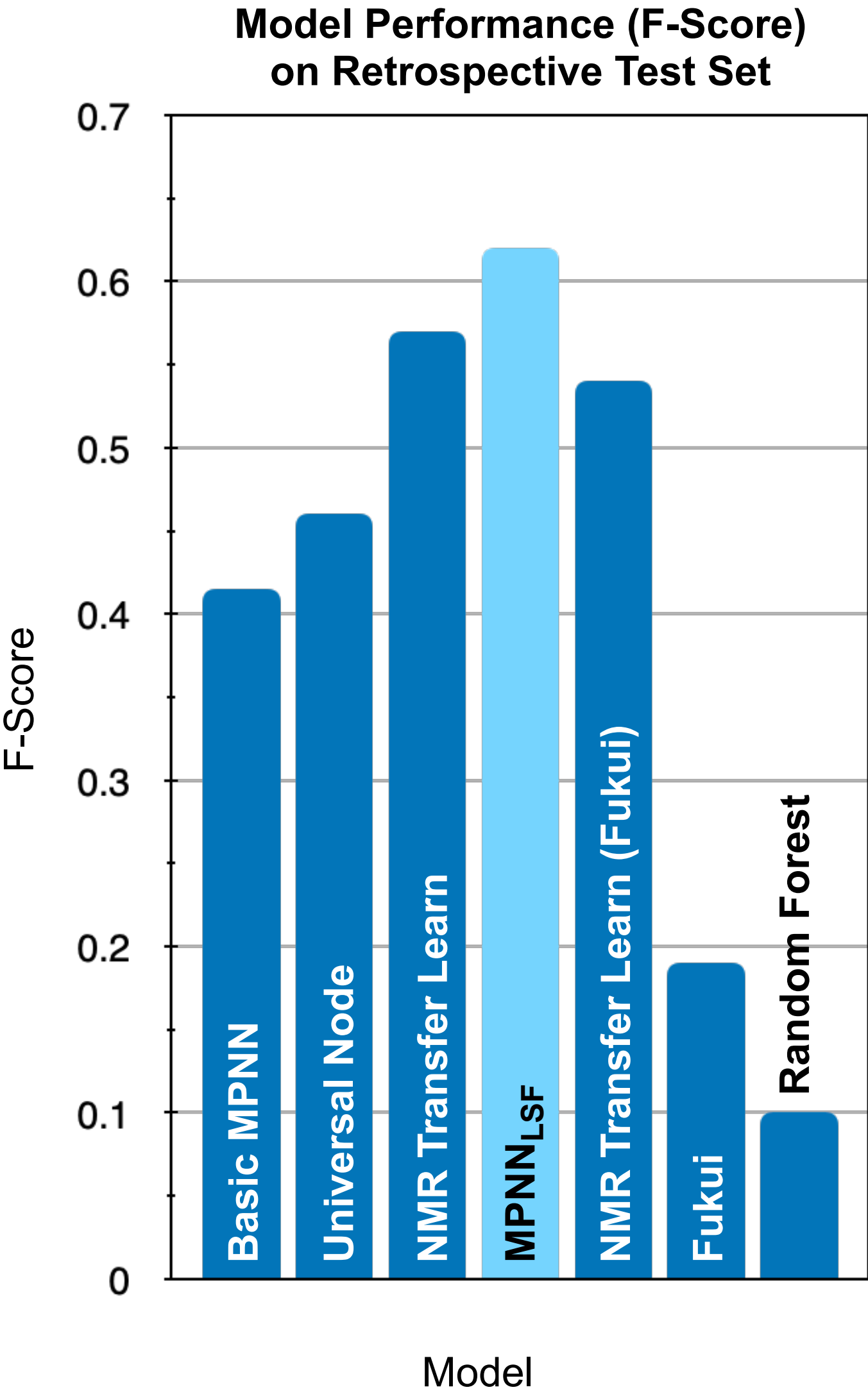
solvent usually **CHCl₃/water**
DMSO/acid mixtures useful for
substrates with limited solubility

DMSO

 increase reactivity for
electron-poor systems

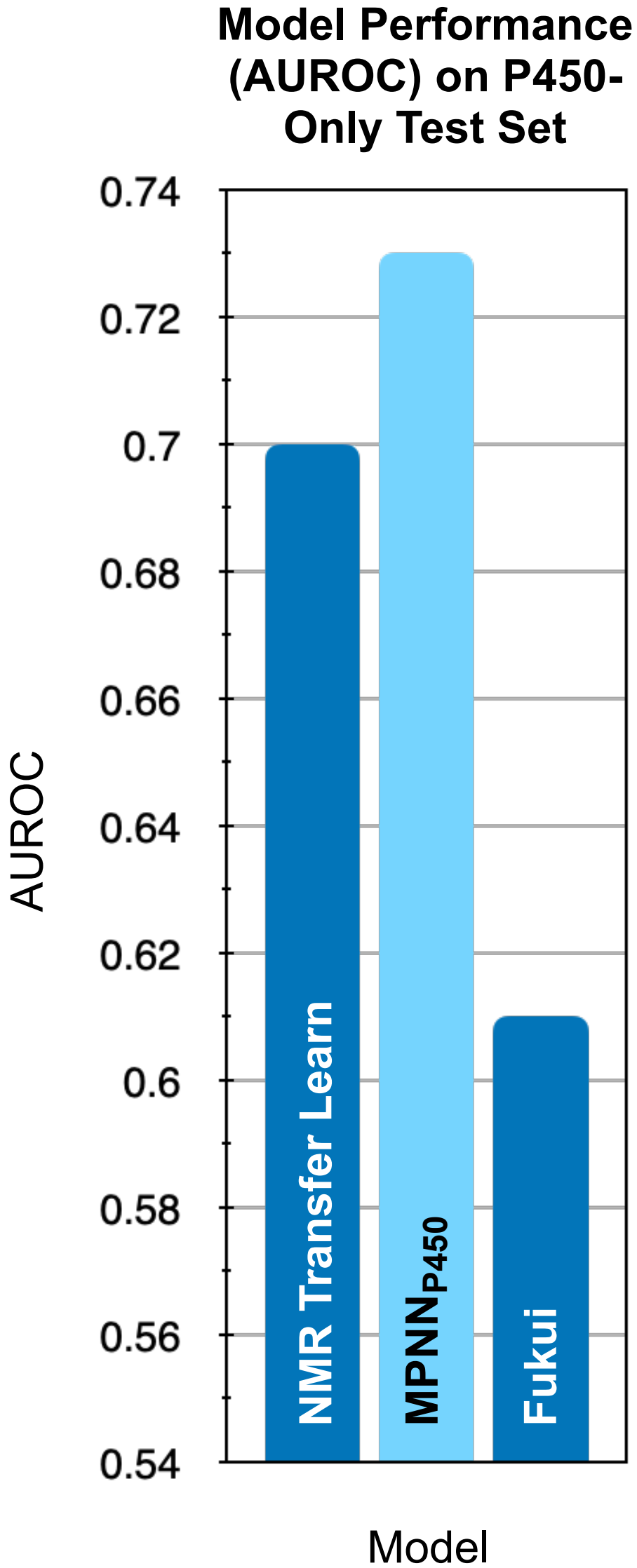
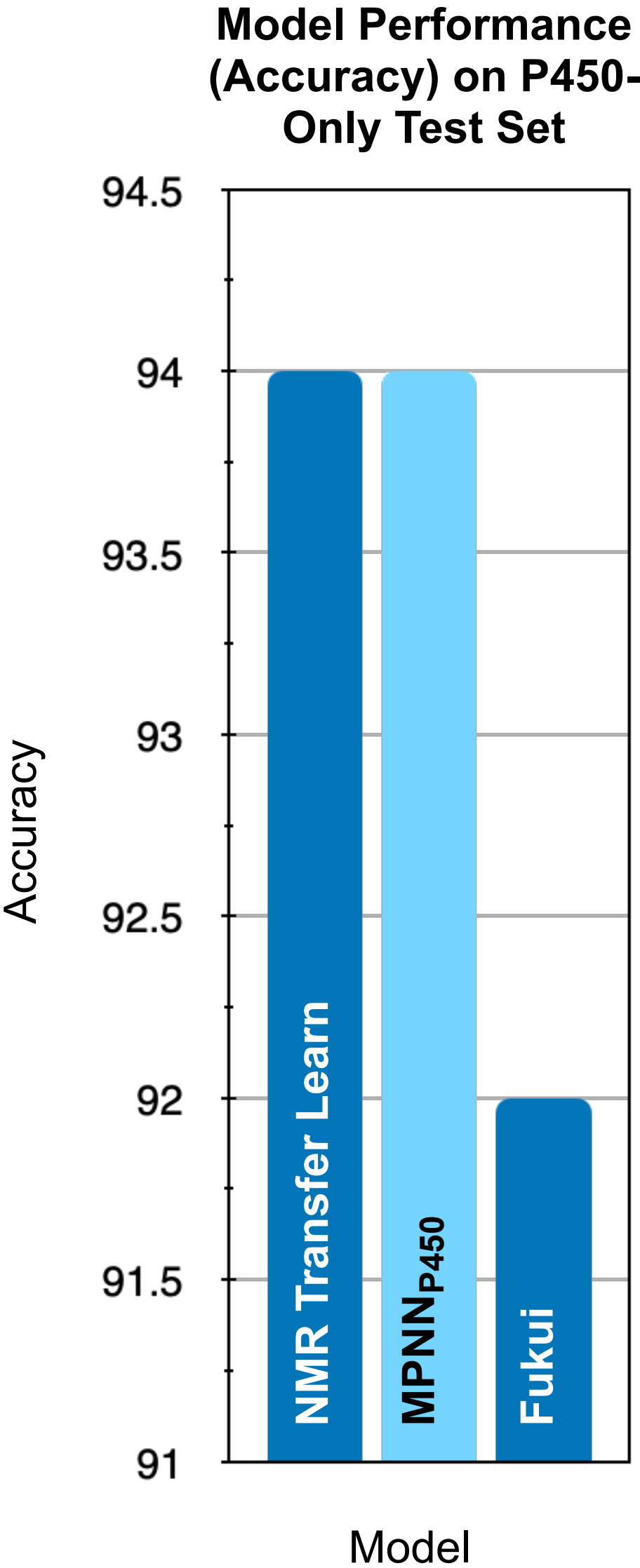
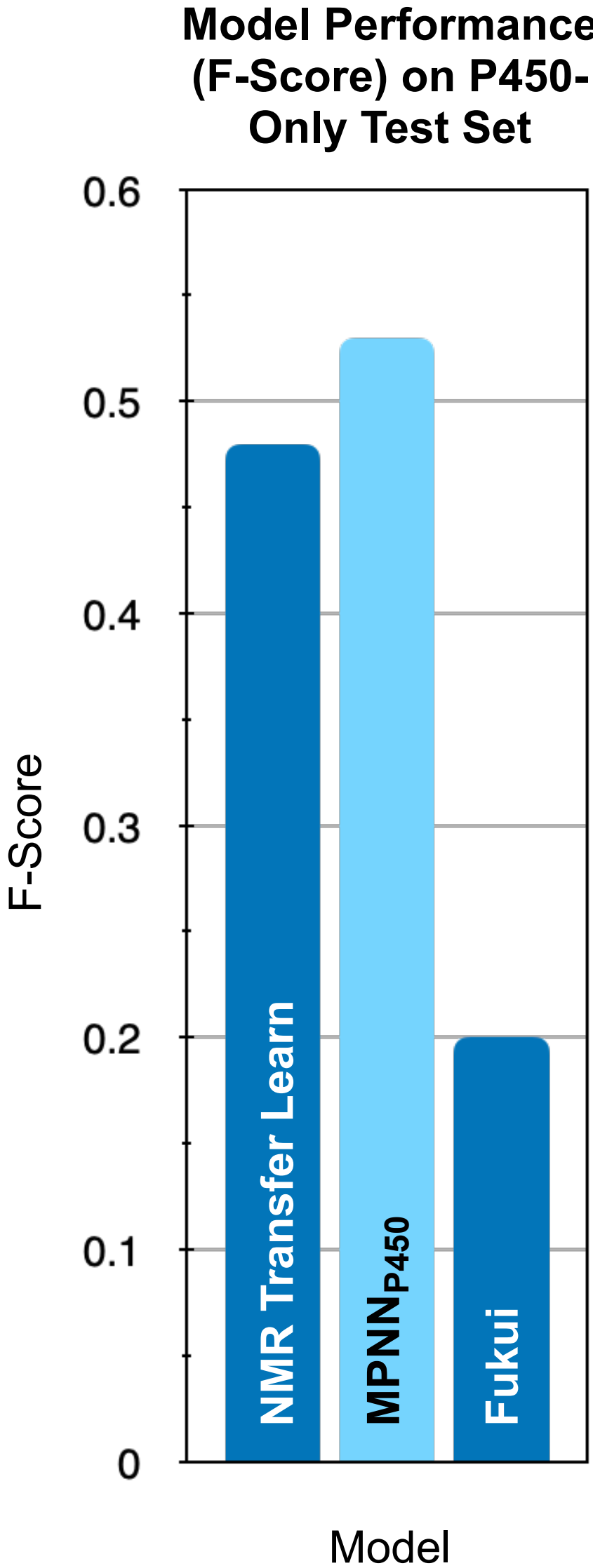
Backup: F-Score, Accuracy, and AUROC

● = Best Model



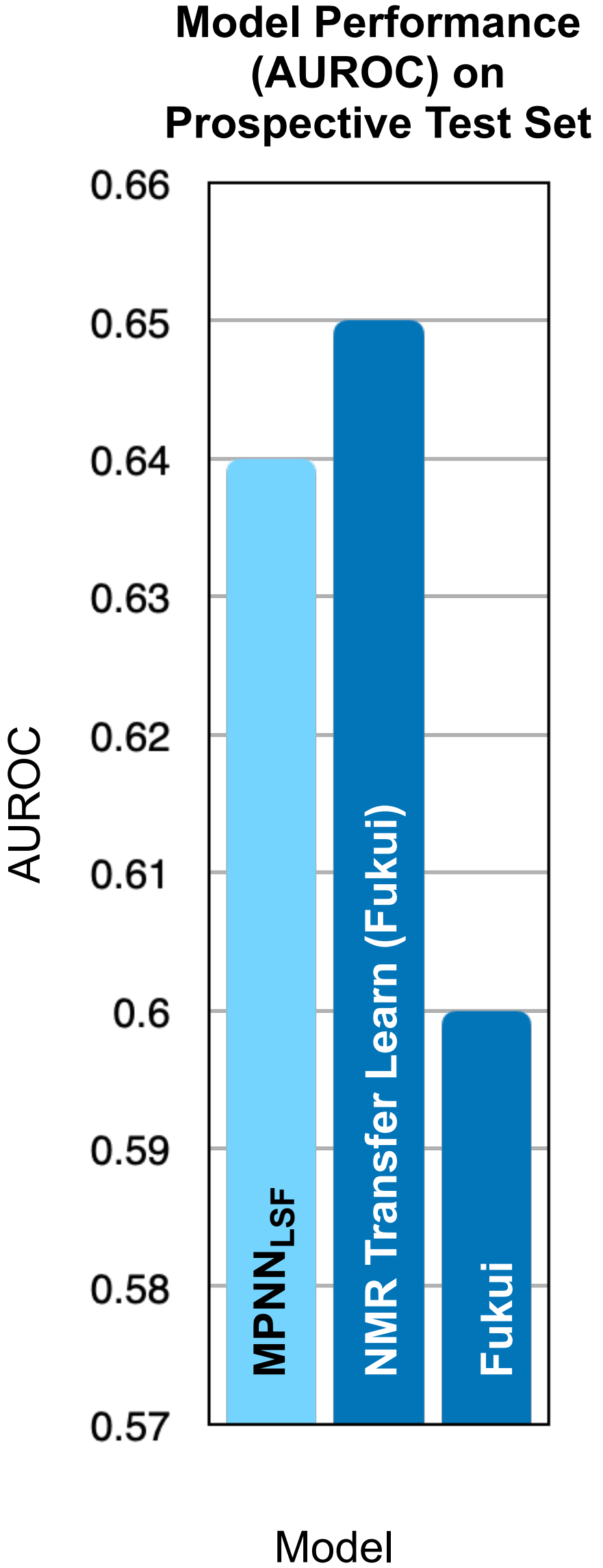
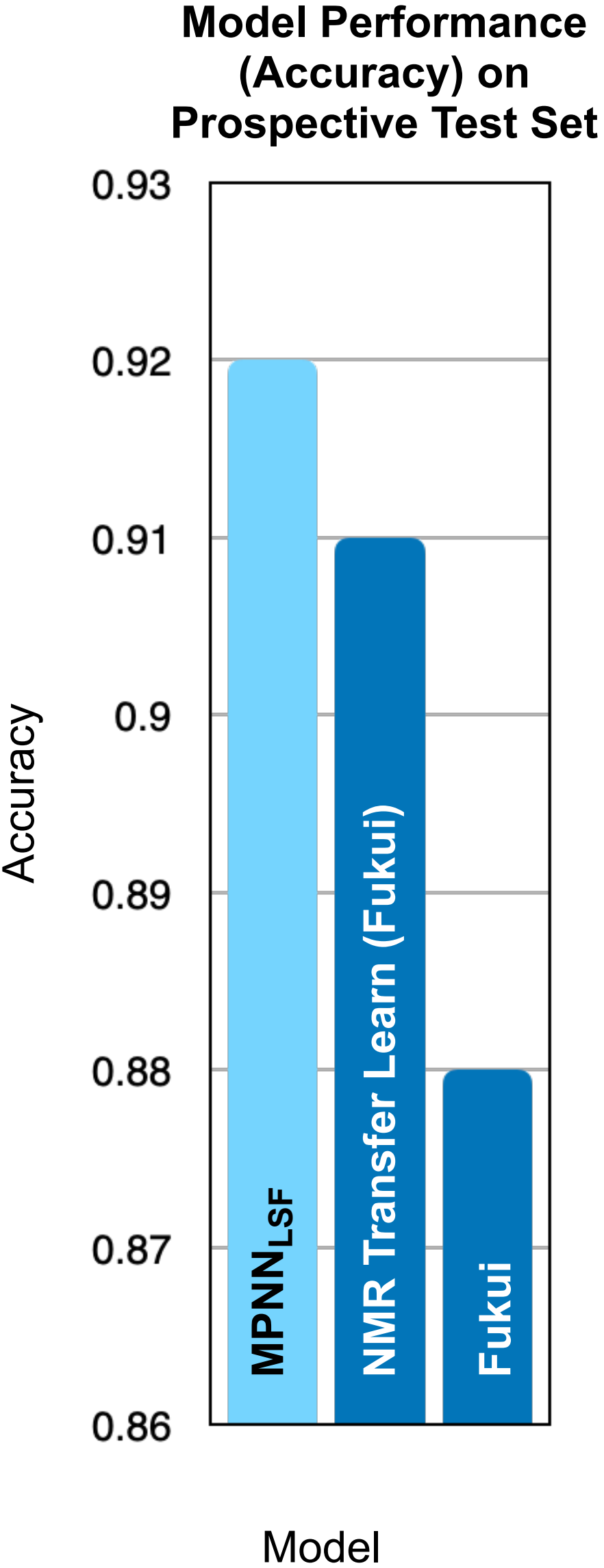
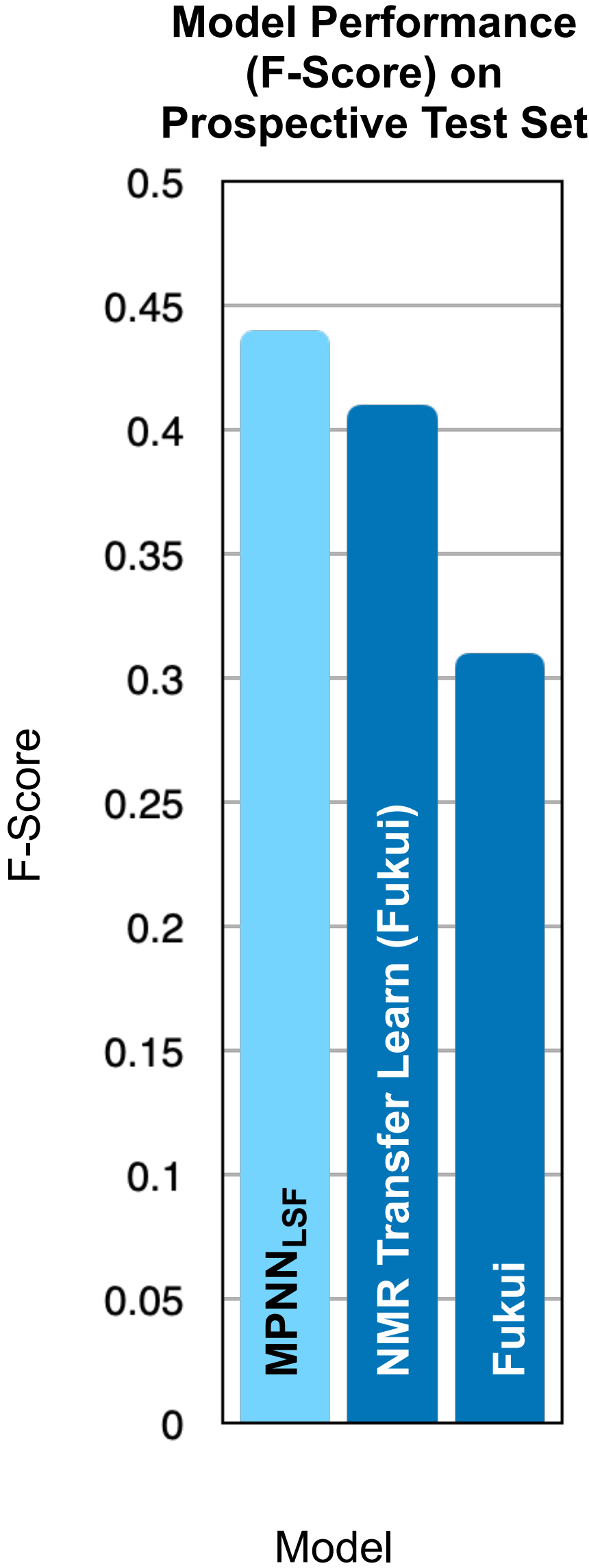
Backup: F-Score, Accuracy, and AUROC

● = Best Model

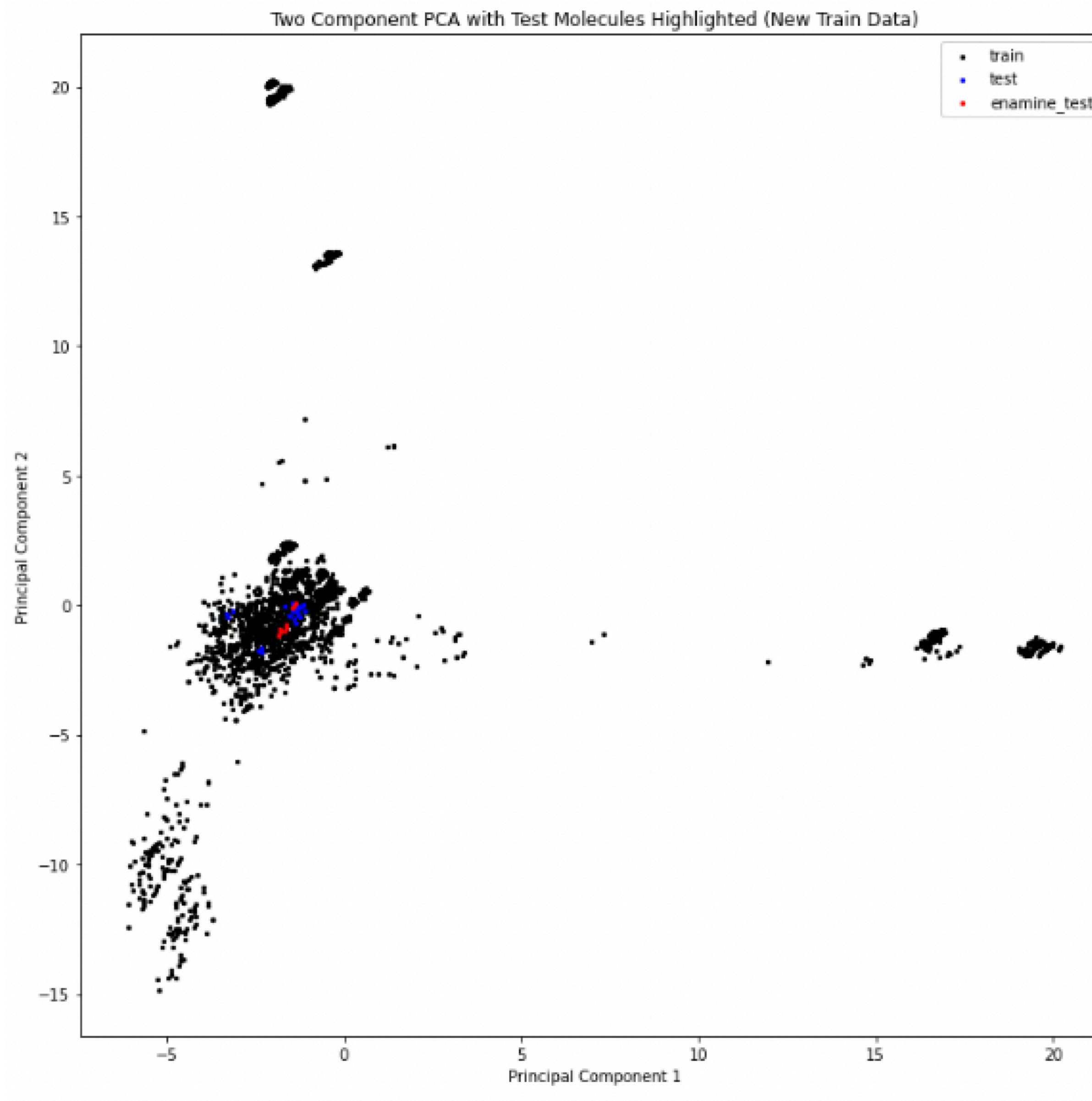


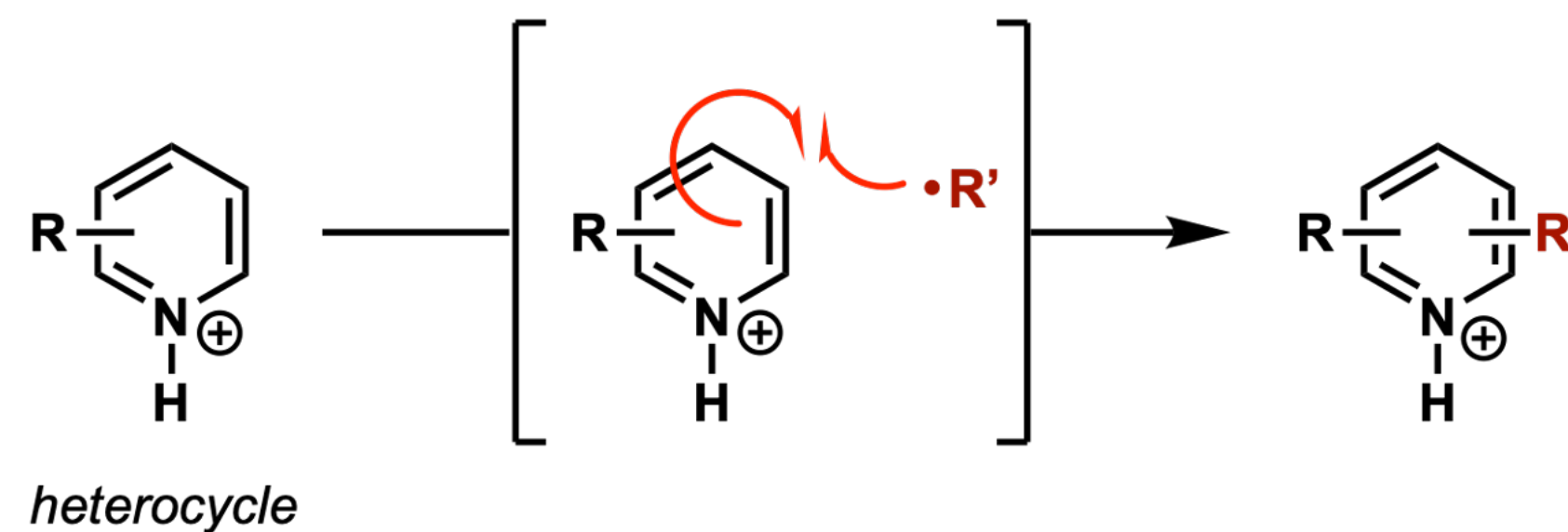
Backup: F-Score, Accuracy, and AUROC

● = Best Model



Backup: PCA of the Dataset Chemical Space





Fukui function-derived predictions

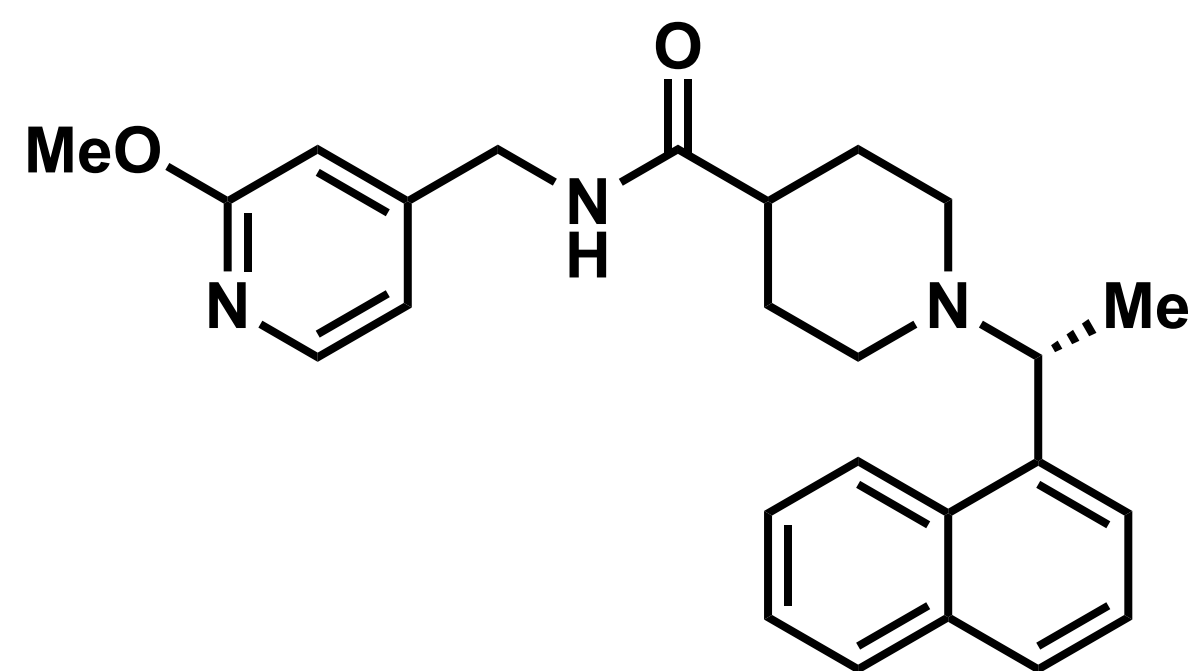
$$F_i(-) = q_i(N - 1) - q_i(N) \quad (\text{electrophilic radicals})$$

$$F_i(0) = \frac{q_i(N - 1) - q_i(N + 1)}{2} \quad (\text{nucleophilic radicals})$$

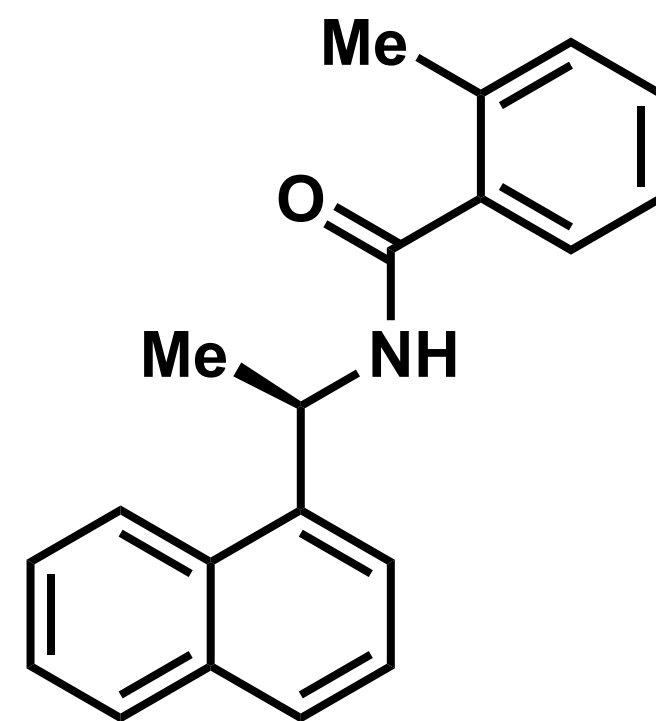
$q_i(N)$ = charge at atom i in a molecule with N electrons.

	F-Score / %	PVV / %	TPR / %	Accuracy / %
aGNN2D	38 (± 5)	56 (± 1)	30 (± 6)	88 (± 1)
aGNN2DQM	39 (± 2)	54 (± 2)	30 (± 3)	87.6 (± 0.3)
aGNN3D	59 (± 3)	62 (± 2)	56 (± 4)	90 (± 1)
aGNN3DQM	60 (± 4)	62 (± 2)	59 (± 6)	90 (± 1)

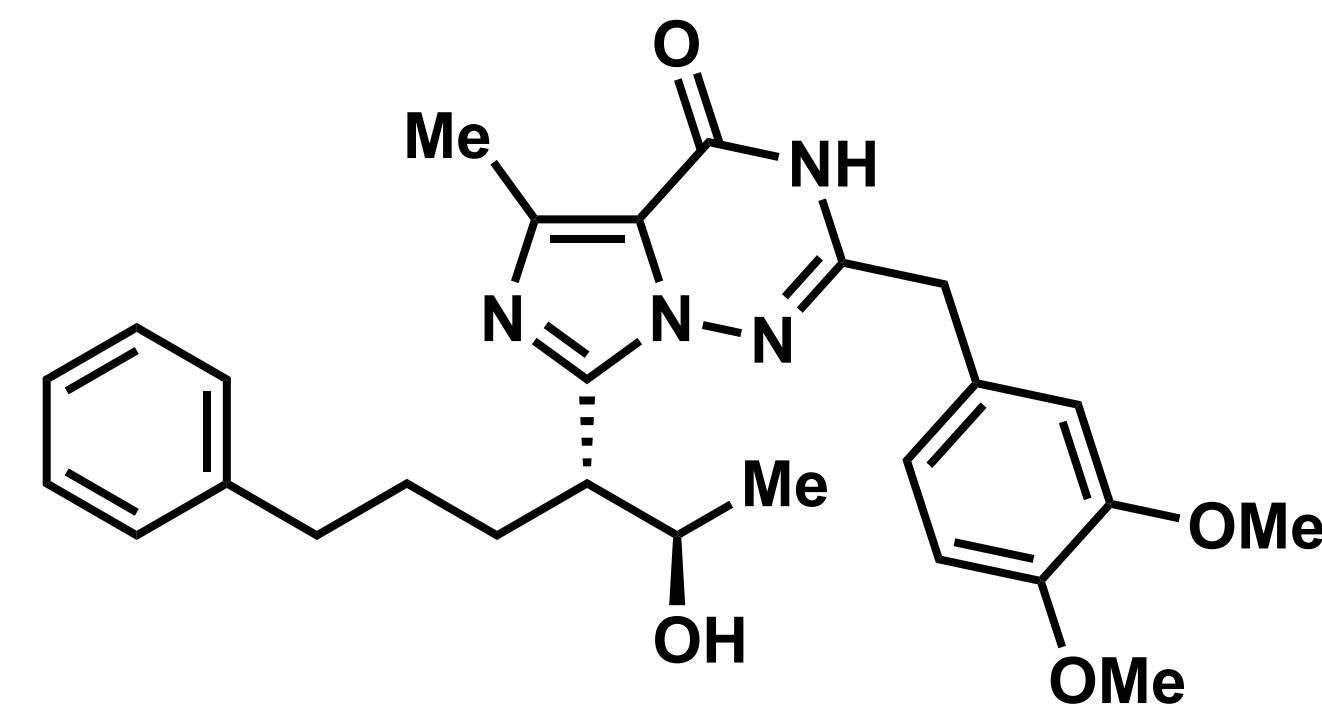
Backup: Representative P450 Test Set Molecules



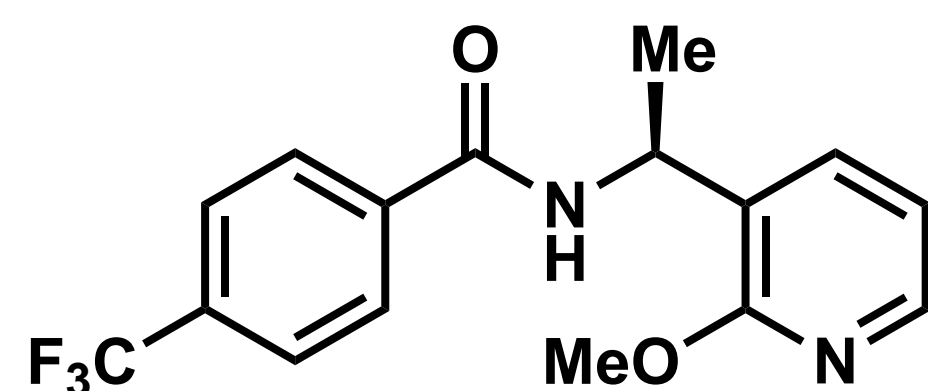
6



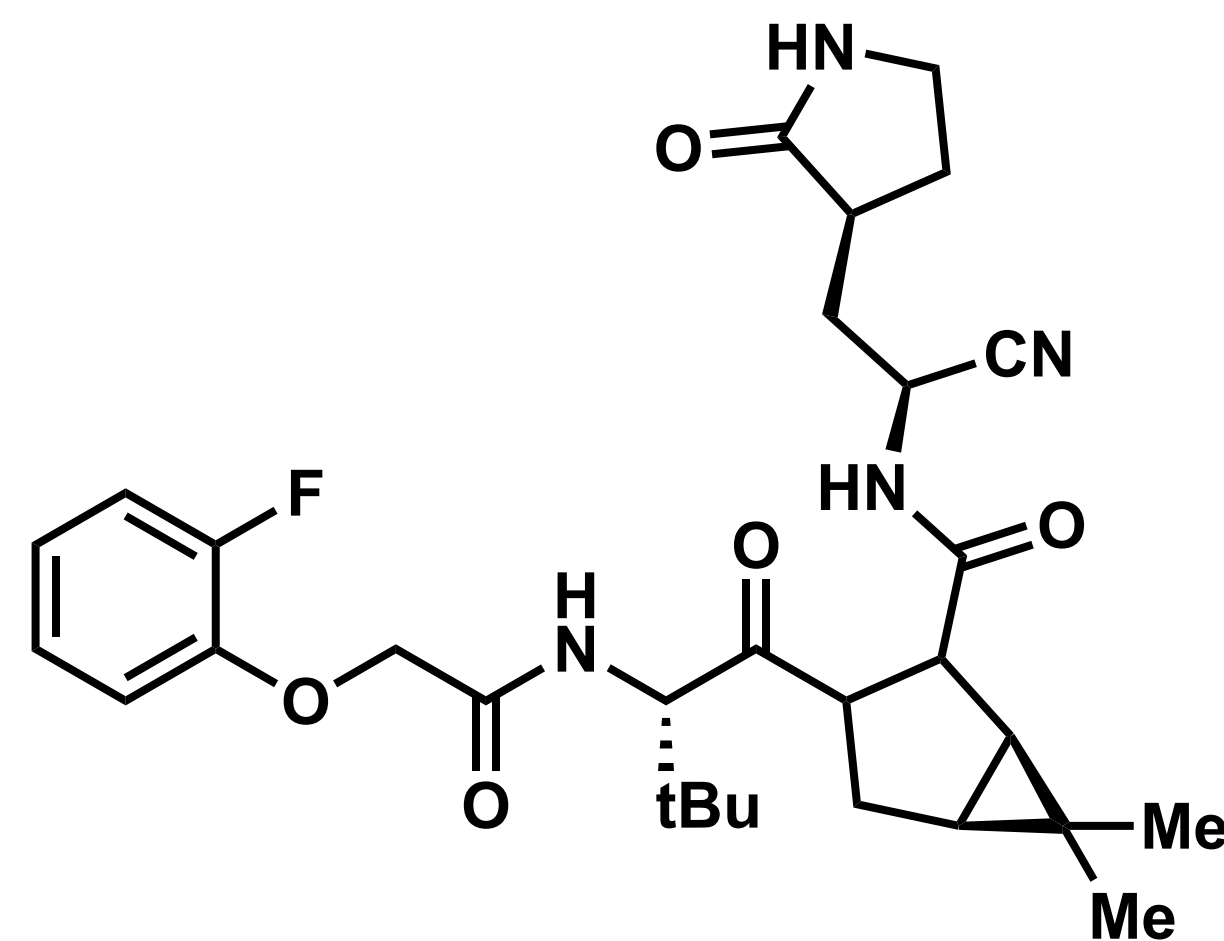
7



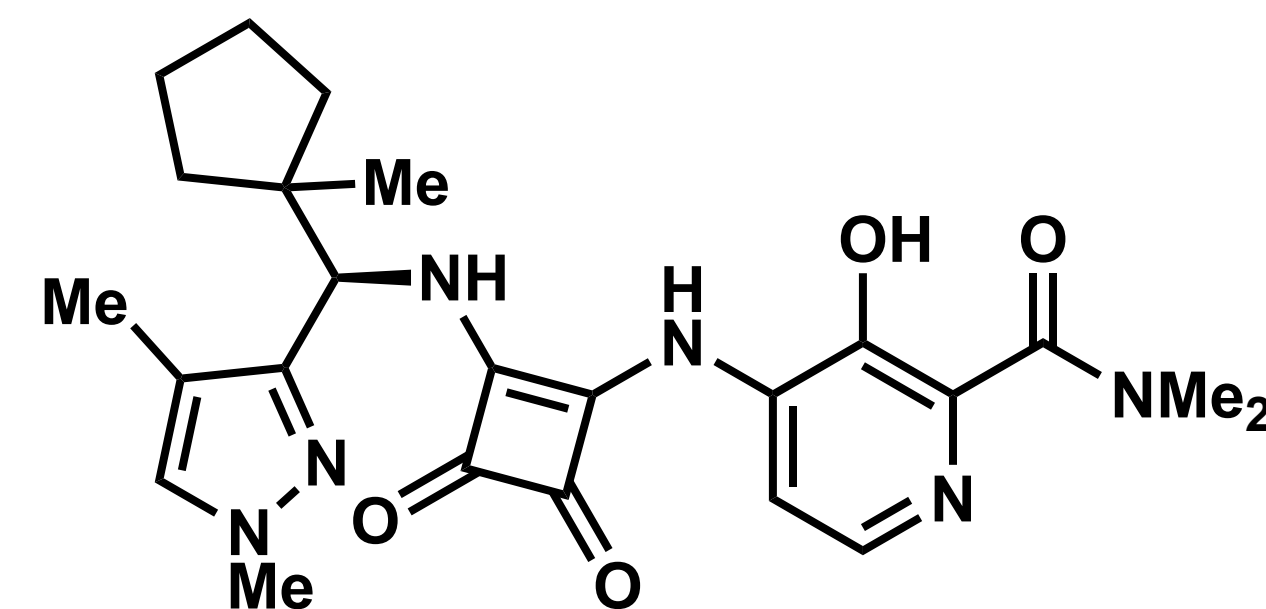
8



9

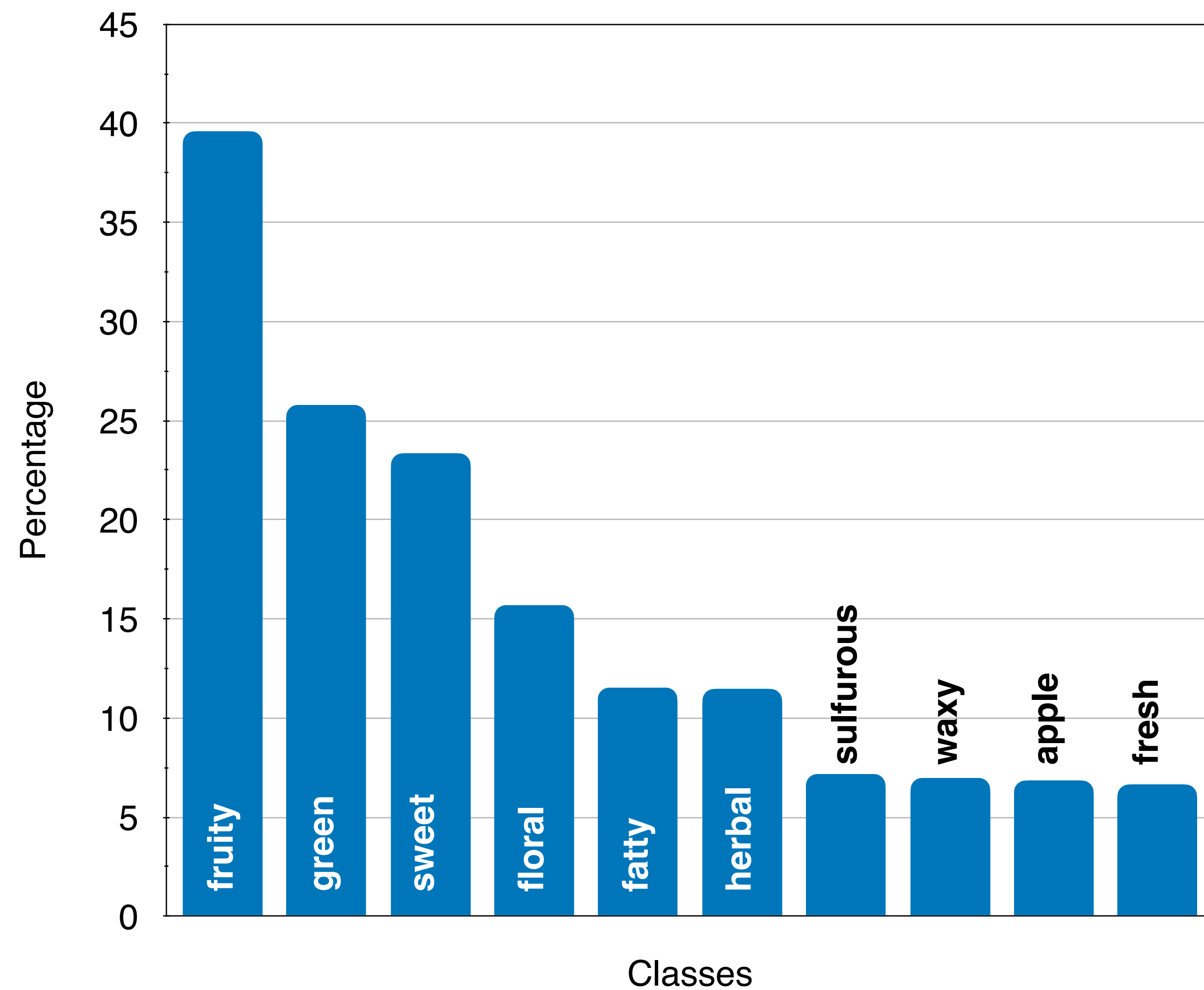


10

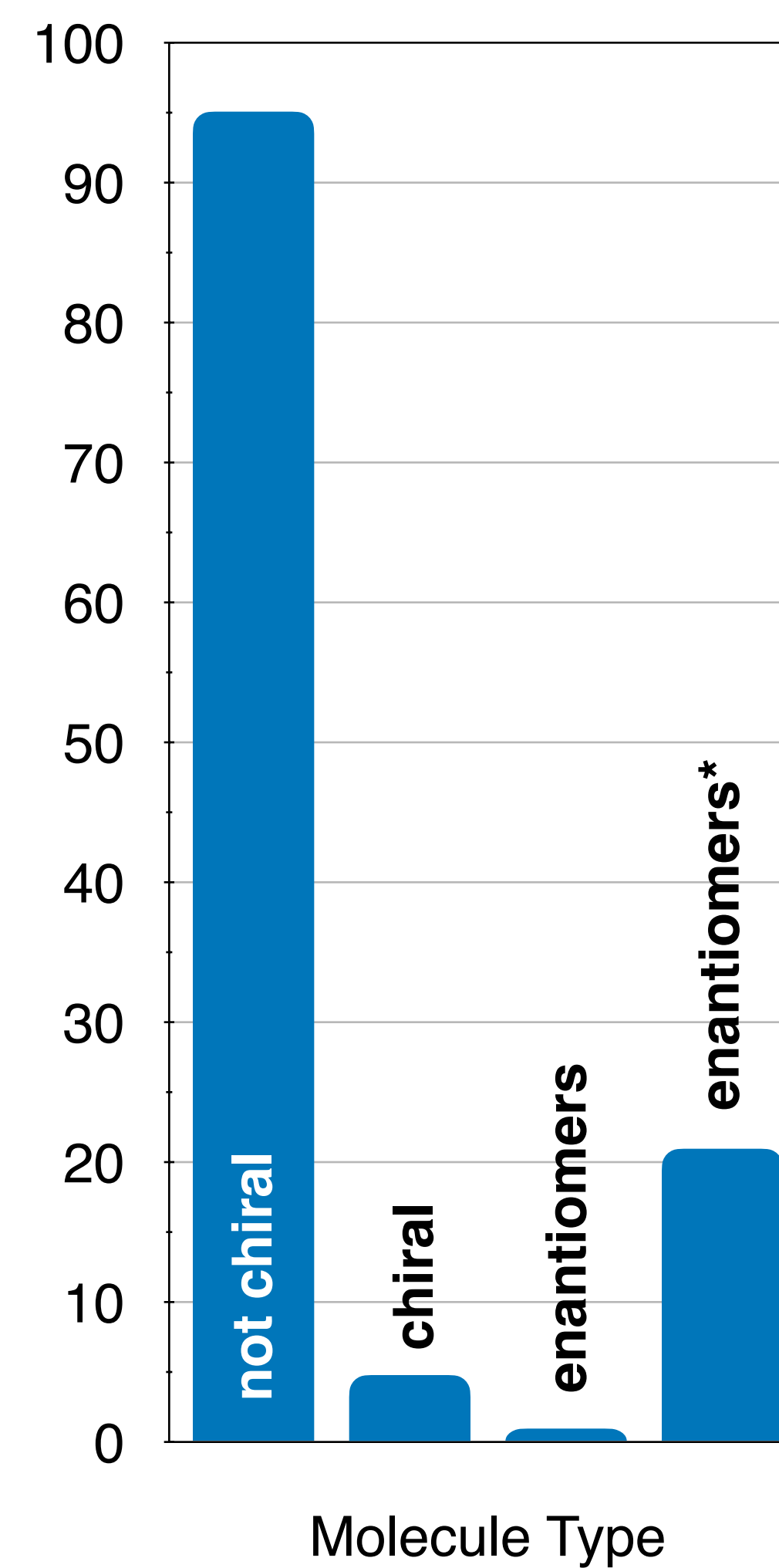


11

Top 10 Odor Classes Distribution

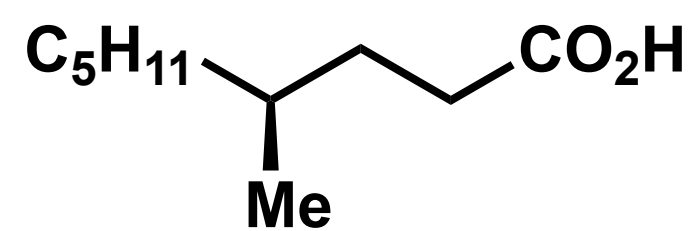


Molecule Distribution



*enantiomers = % enantiomers in all chiral molecules

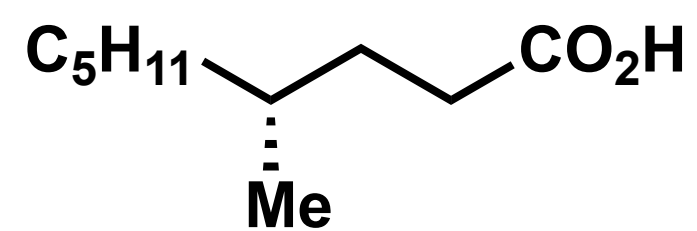
Enantiomeric Pairs Predictions: Similar Olfactive Notes



13

animal

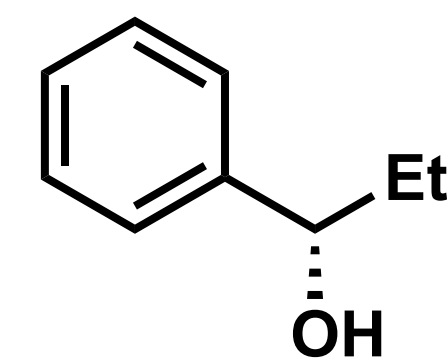
(—)



14

animal

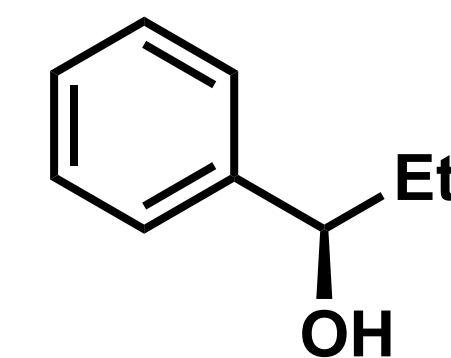
(—)



27

floral

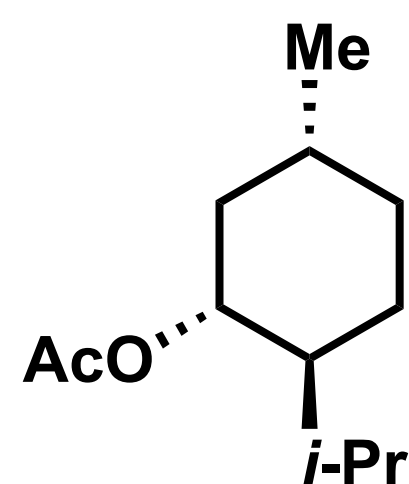
(—)



28

floral

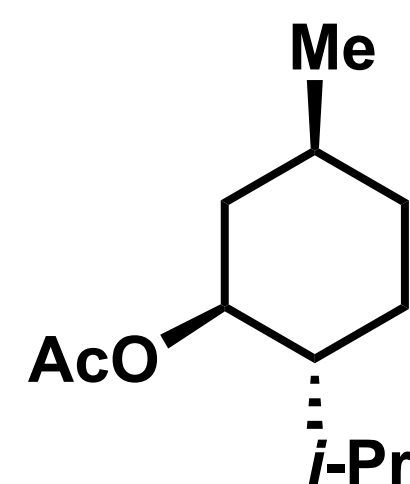
(—)



15

fresh, fruity, mint

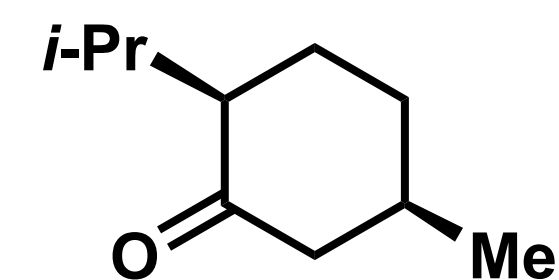
(fruity, mint)



16

fresh, fruity, mint

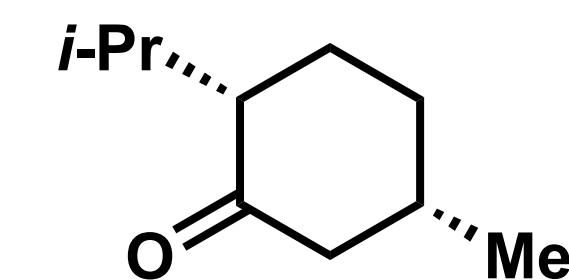
(fruity, mint)



(*R*)-isomethone (17)

mint

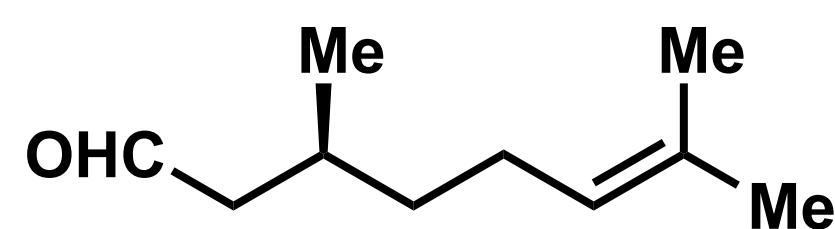
(mint)



(*S*)-isomethone (18)

mint

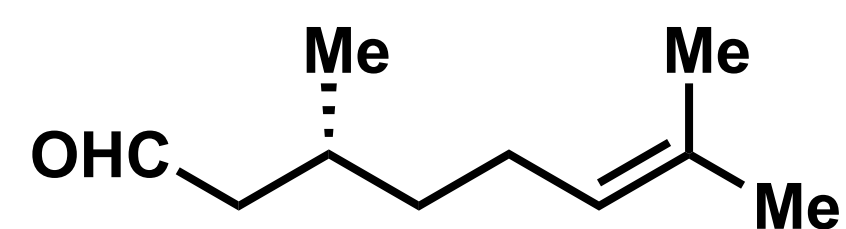
(mint)



(*S*)-citronellal (25)

citrus, fresh, herbal

(fresh, herbal)



(*R*)-citronellal (26)

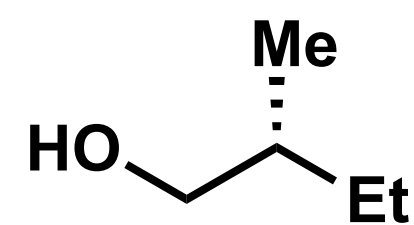
citrus, fresh, herbal

(herbal)

● = correctly identified similarity /
dissimilarity in olfactive notes

● = incorrectly identified similarity /
dissimilarity in olfactive notes

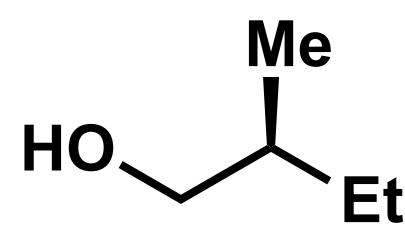
Enantiomeric Pairs Predictions: Dissimilar Olfactive Notes



19

fatty, fermented

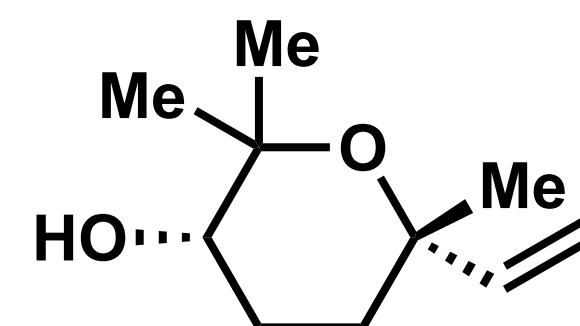
(cheesy*)



20

ethereal, fresh

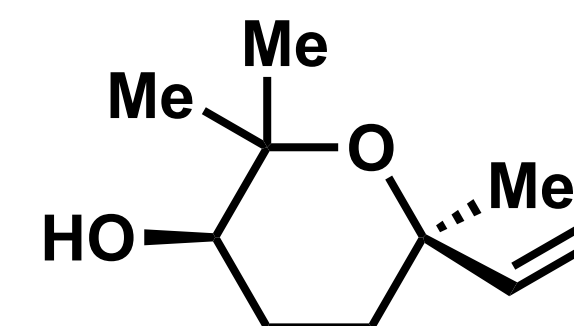
(ethereal)



33

dairy, floral, sweet

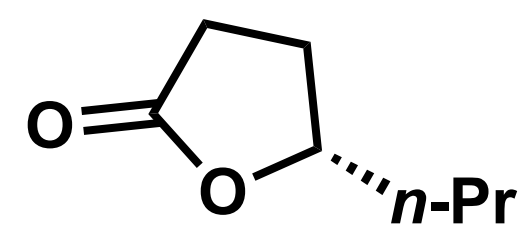
(—)



34

earthy

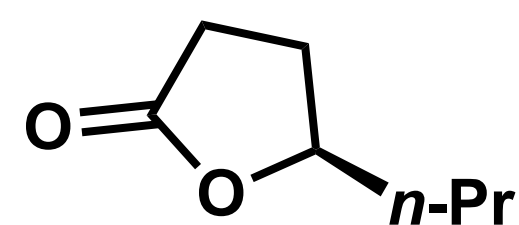
(—)



29

coconut, fruity, sweet

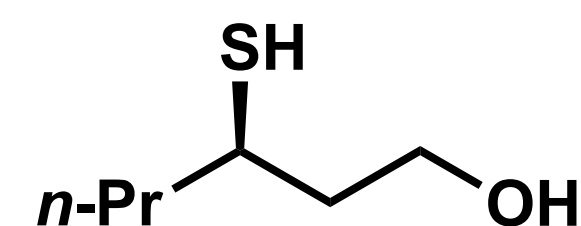
(—)



30

grassy, spicy,
sweet, vanilla

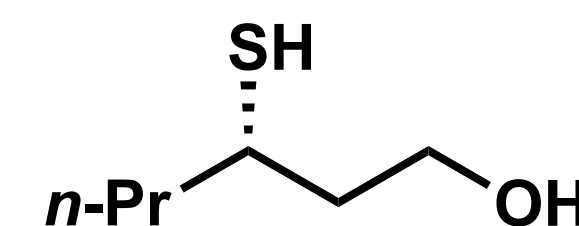
(—)



21

herbal, sulfurous

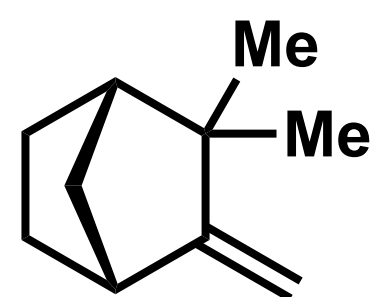
(sulfurous)



22

blackcurrant, fruity,
sweet, topical, woody

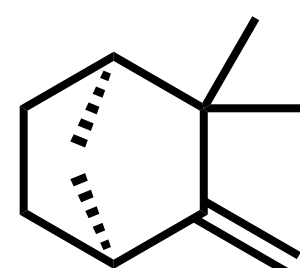
(sweet, fruity)



(*R*)-camphene (**31**)

balsamic, medicinal

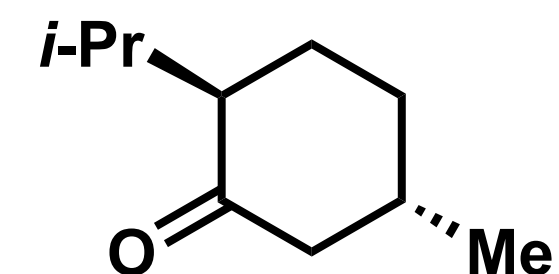
(—)



(*S*)-camphene (**32**)

camphoreous, pine

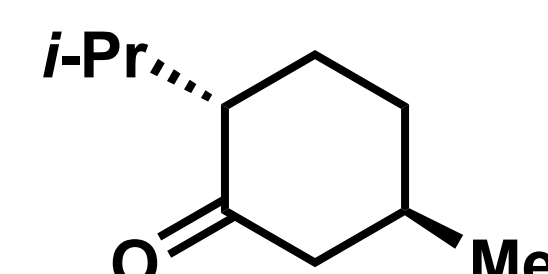
(camphoreous)



(*R*)-methone (**23**)

fresh, musty

(—)



(*S*)-methone (**24**)

camphoreous, fresh

(camphoreous)

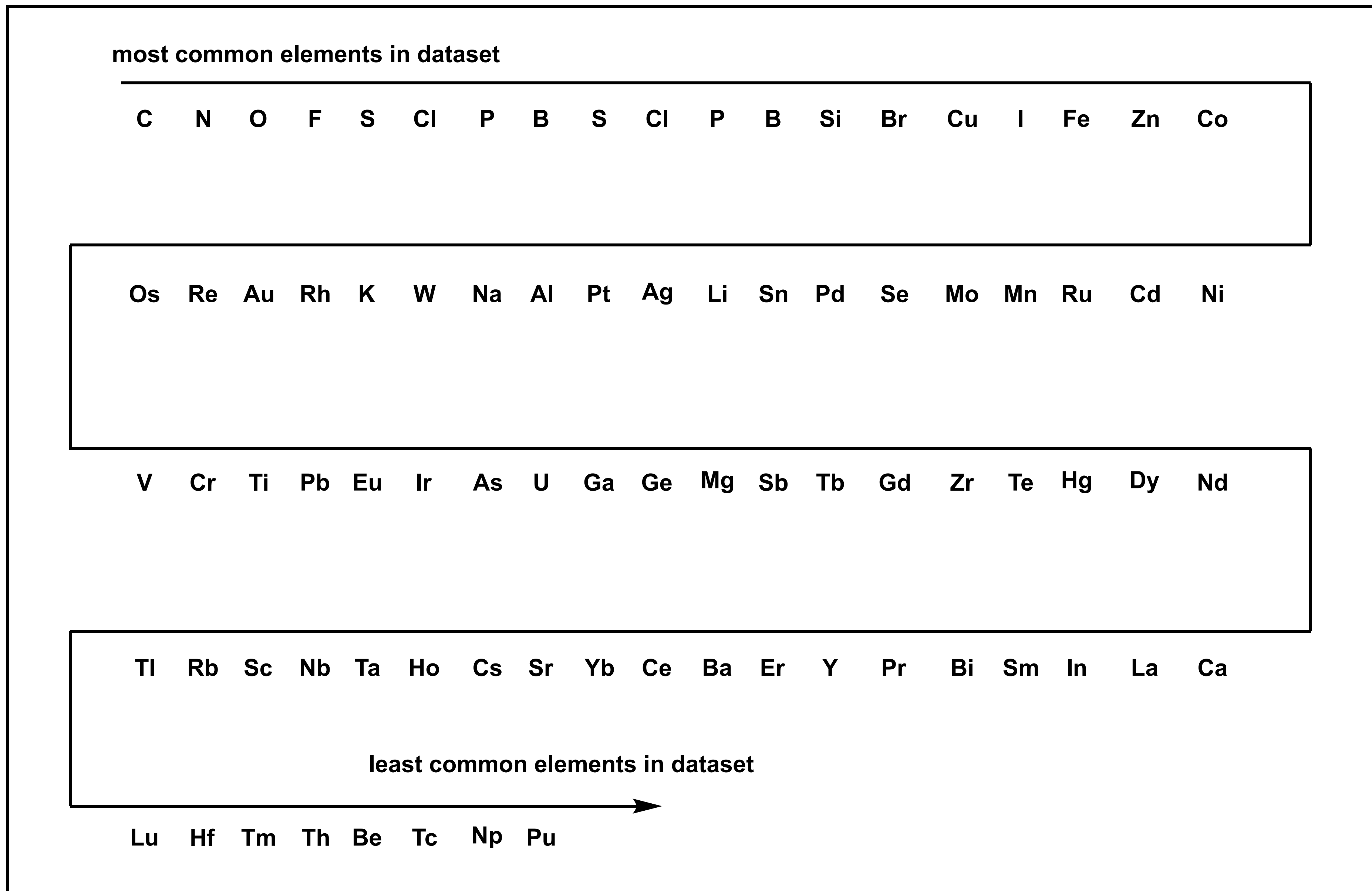
Model	Test Set MSE
Small MPNN	3.17
Large MPNN	2.93

Mean Squared Error (MSE) of total loss (bond distance loss + bond angles loss) on crystal structure data for a variety of message passing neural networks (MPNNs). Test set consisted of unseen molecules.

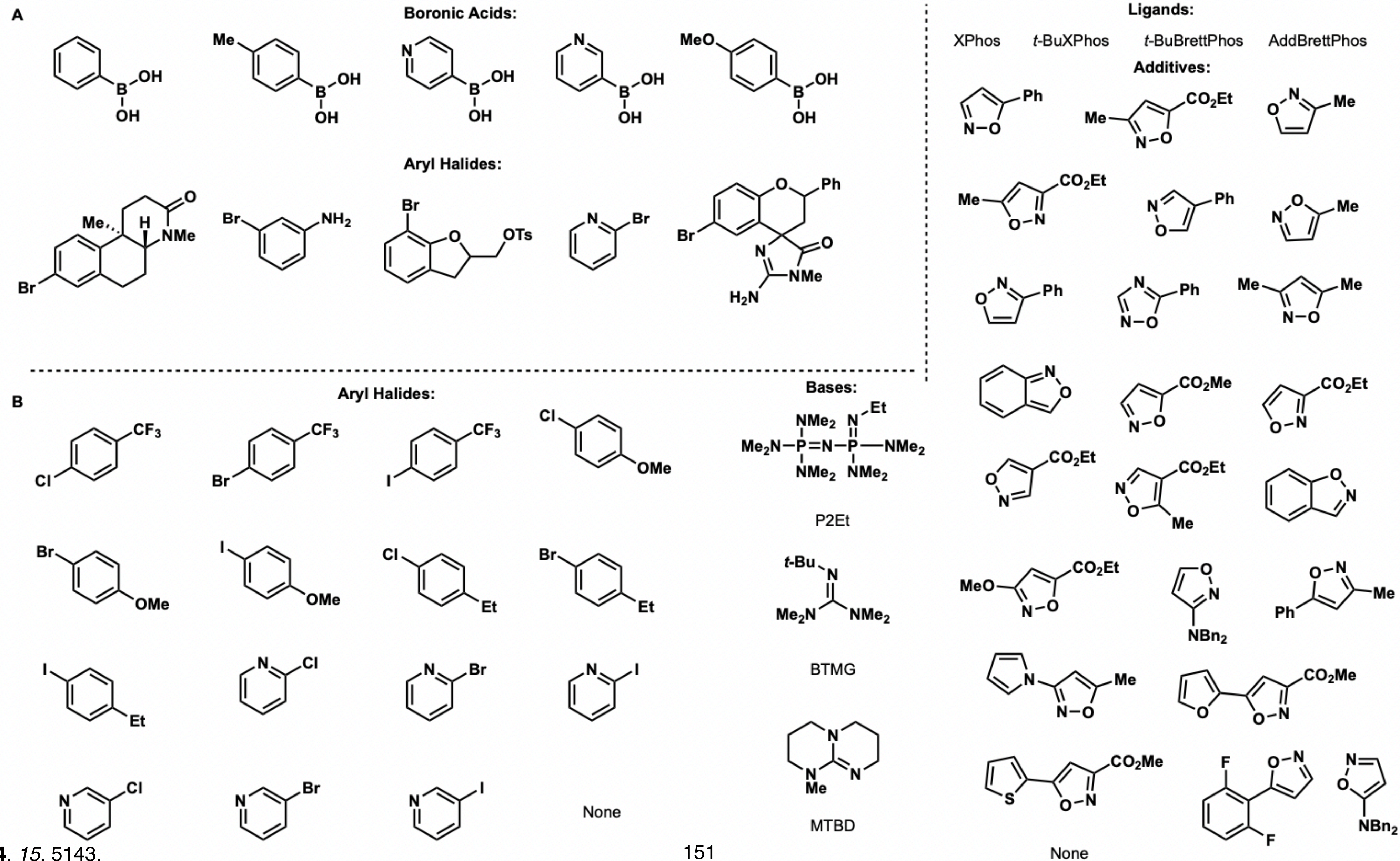
Compound	True Toxicity (log(mol kg⁻¹))	Crystal-Tox Predicted Toxicity (log(mol kg⁻¹))	Oloren ChemEngine Predicted Toxicity (log(mol kg⁻¹))
water	-0.70	1.53	1.98
sucrose	1.06	1.01	1.48
glucose	0.84	1.25	1.77
monosodium glutamate	1.00	1.66	2.10
THC	2.39	2.88	2.53
CBD	2.51	2.62	2.41
aconitine	6.90	3.84	3.38
epibatidine	7.43	2.88	2.93
MDMA	3.08	2.59	2.55
cocaine	3.50	2.09	2.67
LSD	4.29	2.65	2.89
heroin	4.23	2.80	3.19

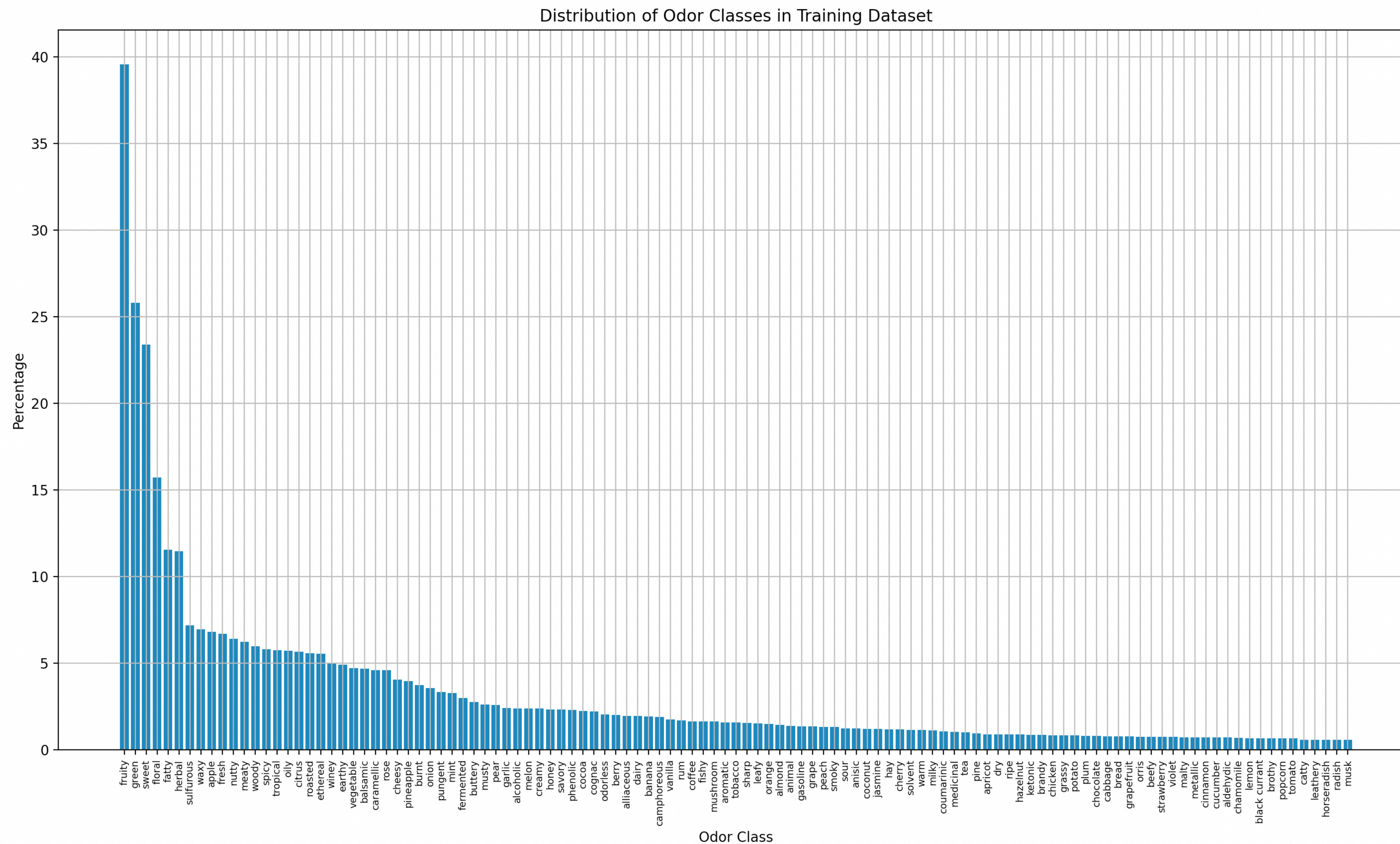
Per-Molecule Set Error Rate (Yields)

Model	Split MAE			
	<i>Halide Set 0</i>	<i>Halide Set 1</i>	<i>Halide Set 2</i>	<i>Halide Set 3</i>
Random Forest	23.6	23.9	22.2	31.0
Gaussian Process	27.3	25.2	21.7	30.9
Adaboost	24.6	23.9	18.7	31.6
Yield-BERT	27.3	25.2	21.7	30.9
GraphRXN	9.5	41.6	30.9	18.7
Crystal-Yield	26.7	14.8	16.3	27.5
	<i>Base 0</i>	<i>Base 1</i>	<i>Base 2</i>	
Random Forest	32.0	32.4	19.9	
Gaussian Process	31.0	34.3	24.8	
Adaboost	27.2	29.5	19.9	
Yield-BERT	23.3	27.4	22.1	
GraphRXN	12.8	27.1	13.8	
Crystal-Yield	13.9	13.0	13.4	
	<i>Ligand 0</i>	<i>Ligand 1</i>	<i>Ligand 2</i>	<i>Ligand 3</i>
Random Forest	27.4	29.0	27.6	29.8
Gaussian Process	39.8	32.2	29.2	30.6
Adaboost	26.8	29.9	25.9	27.2
Yield-BERT	20.4	24.0	25.8	27.0
GraphRXN	9.7	17.6	12.7	15.2
Crystal-Yield	24.5	23.4	10.4	14.5
Crystal-Yield ^a	17.1	12.2	6.5	10.8
	<i>Additive Set 0</i>	<i>Additive Set 1</i>	<i>Additive Set 2</i>	<i>Additive Set 3</i>
Random Forest	34.0	31.3	26.7	29.4
Gaussian Process	32.7	29.0	24.5	27.9
Adaboost	29.0	27.3	26.7	27.5
Yield-BERT	25.2	22.9	22.8	25.3
GraphRXN	16.7	15.2	22.8	15.4
Crystal-Yield	15.6	16.6	17.2	15.5

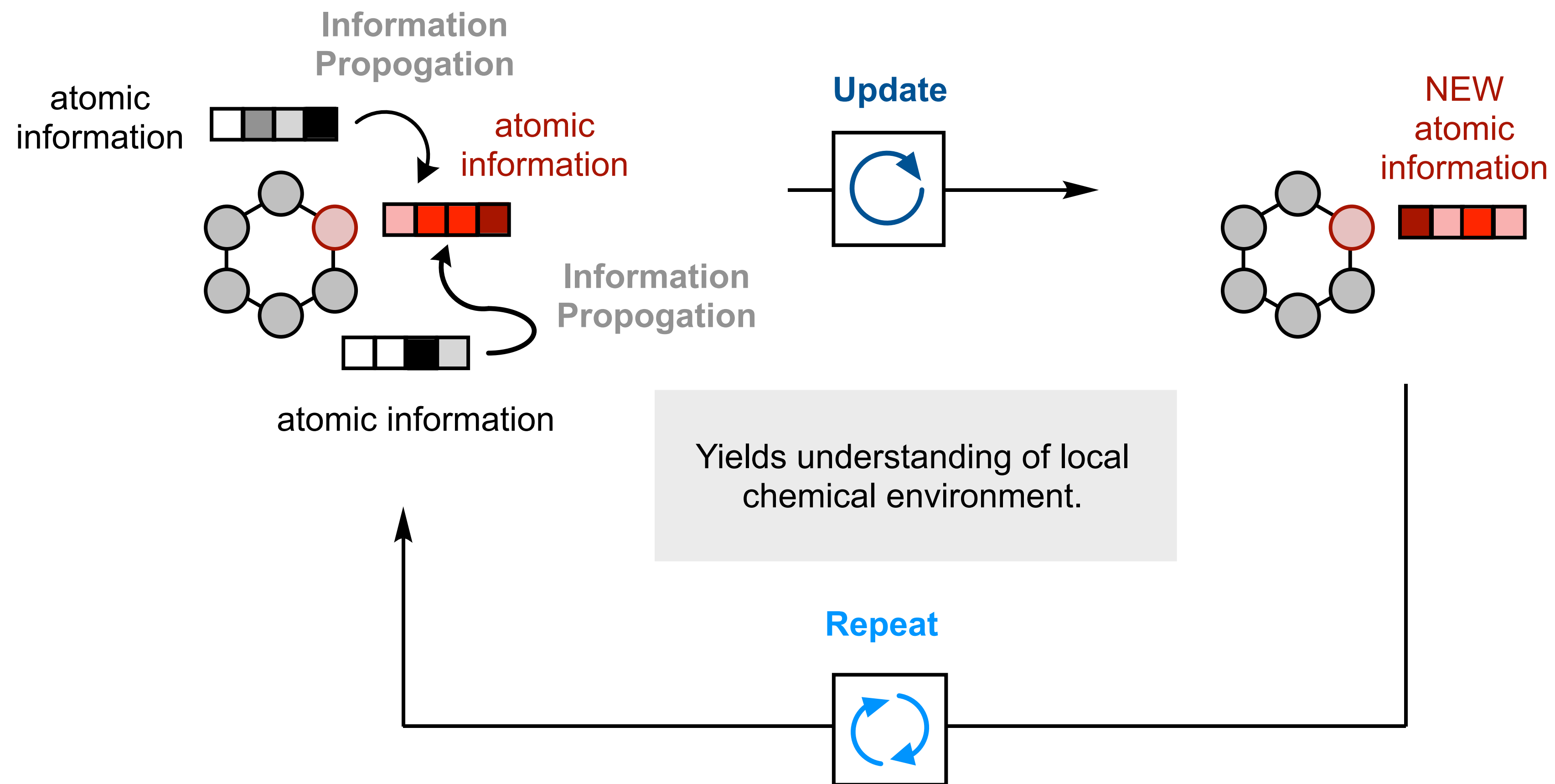


Selected Molecules in Yield Datasets

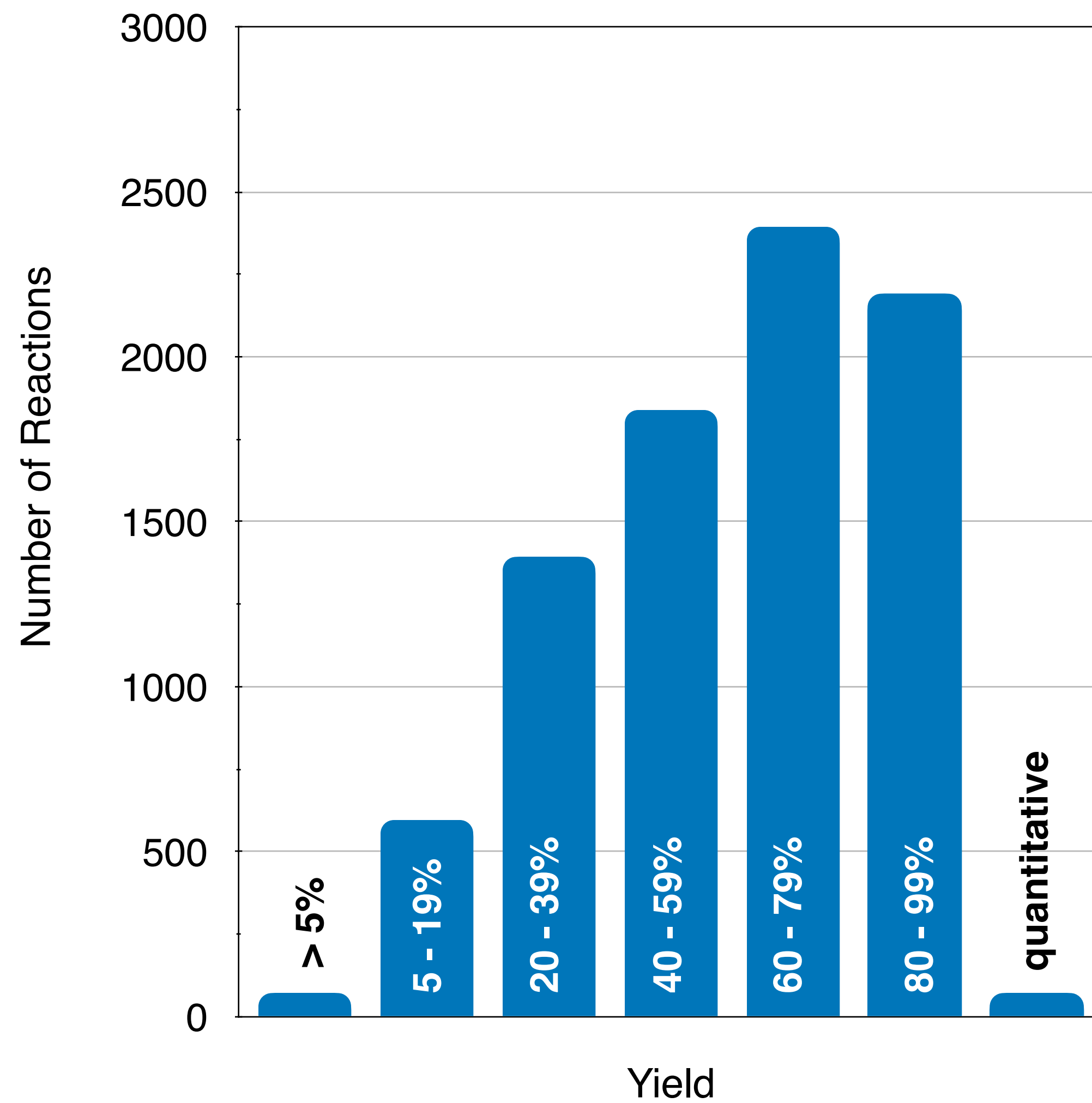




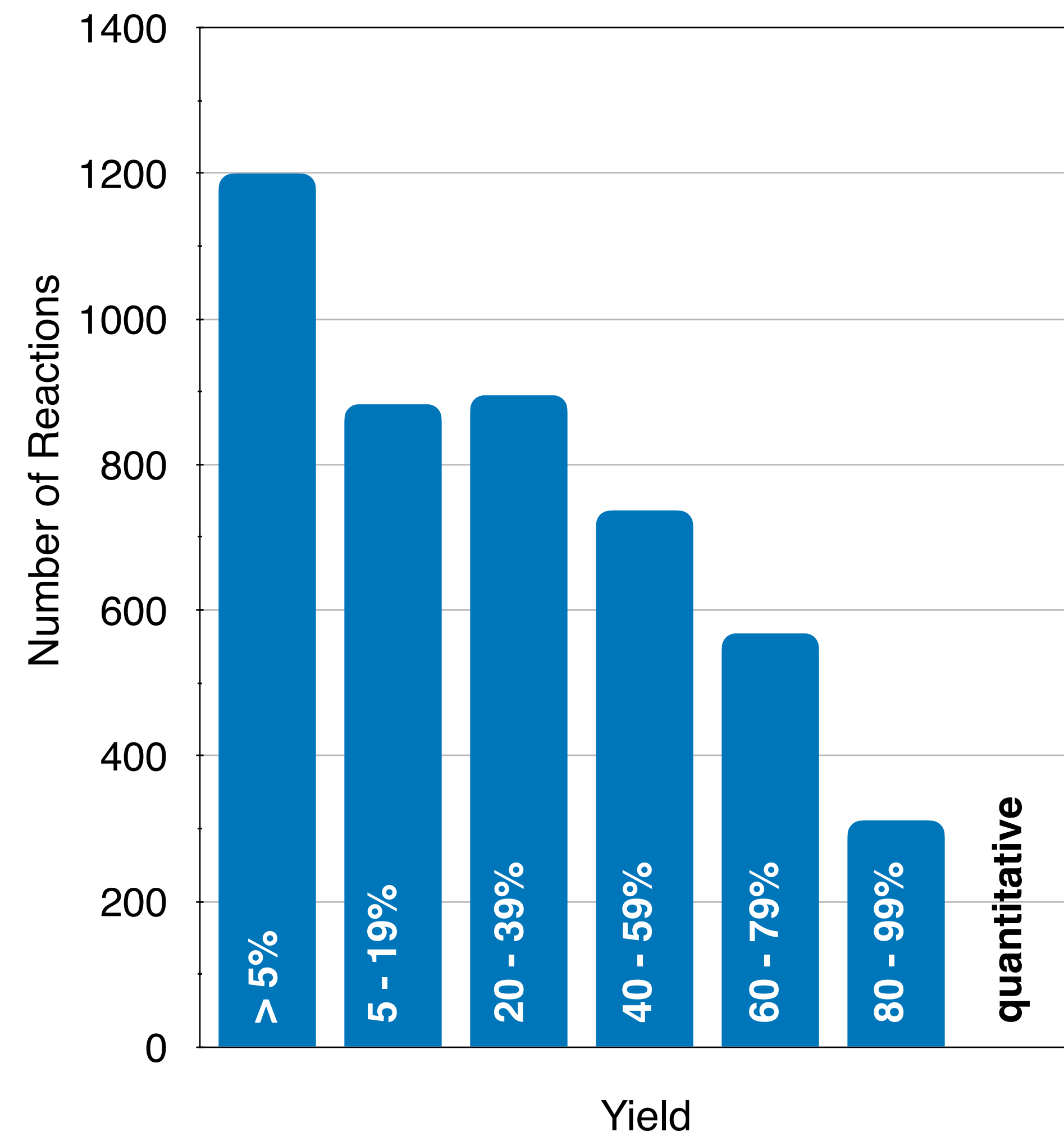
How Does the MPNN Work?



Suzuki USPTO Yield Distribution



Buchwald-Hartwig HTE Yield Distribution



The Big Idea

