SUPPLEMENTARY FILE

Unveiling Multifunctional Inhibitors: Holistic Spectral, Electronic and Molecular

Characterization Coupled with Biological Profiling of Substituted Pyridine Derivatives

Against LD Transpeptidase, Heme Oxygenase and PPAR Gamma

Shaik Yasmin Begum^a, Predhanekar Mohamed Imran^{a,*}, Attar Kubaib^{a,*}, Mohamed Taha Yassin^b, Fatimah O. Al-Otibi^b, M. Selvakumaran^a, A. Aathif Basha^c,

S. Sulthanudeen^a

^aDepartment of Chemistry, Islamiah College (Autonomous), Vaniyambadi - 635752, Tamilnadu, India. (Affiliated to Thiruvalluvar University, Serkkadu, Vellore – 632115, Tamilnadu, India) ^bDepartment of Botany and Microbiology, College of Science, King Saud University, P.O. Box 2455, Riyadh 11451, Saudi Arabia

^cDepartment of Physics, Islamiah College (Autonomous), Vaniyambadi - 635752, Tamilnadu, India. (Affiliated to Thiruvalluvar University, Serkkadu, Vellore – 632115, Tamilnadu, India)

Figure S1. Chemical structure of 3-Phenyl-1-(Pyridine-4-Carbonyl)-1H-Pyrazole-4-Carboxylic Acid (PPP), 3-(4-Acetophenyl-)1-(Pyridine-4-Carbonyl)-1H-Pyrazole-4-Carboxylic Acid (APP) and 3-(4-Methylphenyl)-1-(Pyridine-4-Carbonyl)-1H-Pyrazole-4-Carboxylic Acid (MPP) derivatives

Figure S2. NMR and mass spectrometry data for all derivatives, obtained through experimental
Figure S3. The binding interactions of compound PPP against LD Transpeptidase (PDB ID: 4JMN), Heme Oxygenase (PDB ID: 1IW0) and PPAR Gamma (PDB ID: 2ZNO)
Figure S4. The binding interactions of compound MPP against LD Transpeptidase (PDB ID: 4JMN), Heme Oxygenase (PDB ID: 1IW0) and PPAR Gamma (PDB ID: 2ZNO)
Table S5. Comprehensive Physicochemical, Medicinal Chemistry, Absorption, Distribution,

Metabolism, Excretion and Toxicity Profiles of Compounds PPP, APP and MPP

*Corresponding authors: imranpkm@gmail.com, attar.kubaib@gmail.com

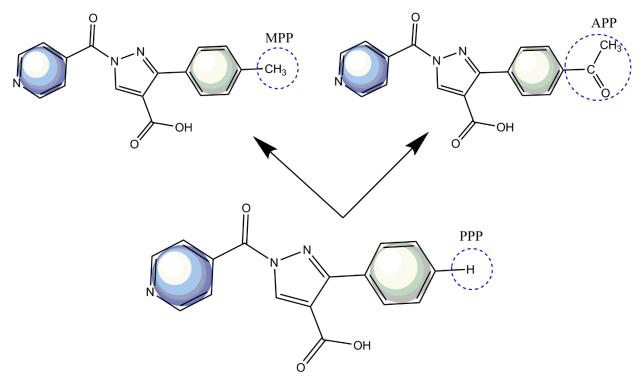
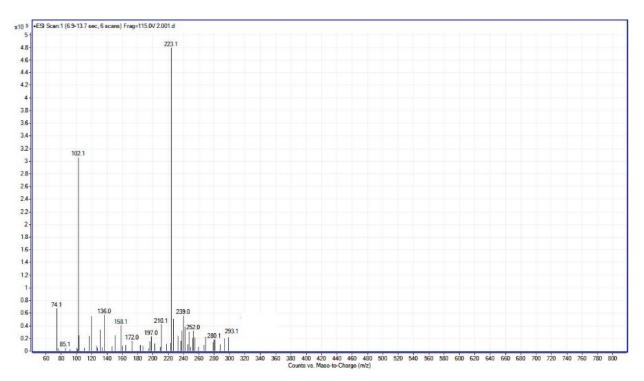
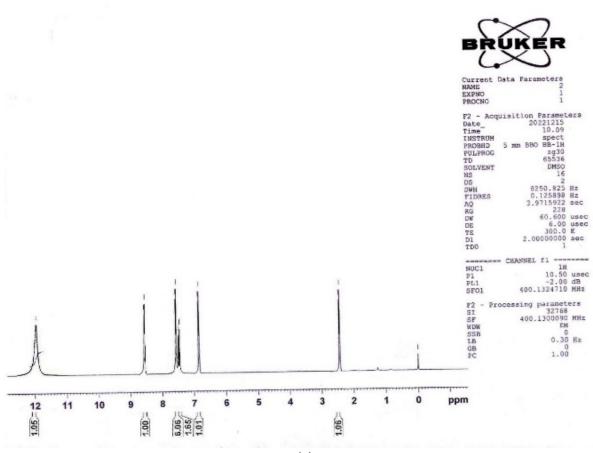
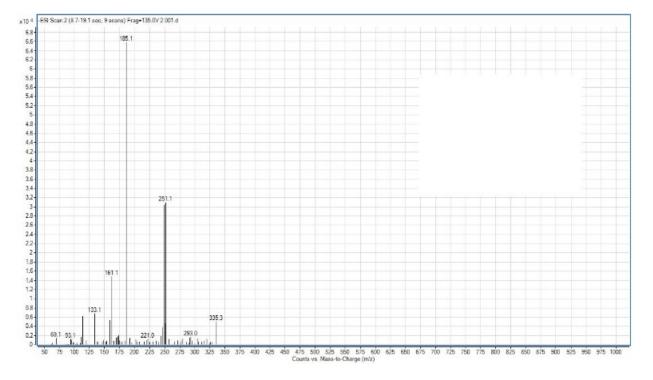


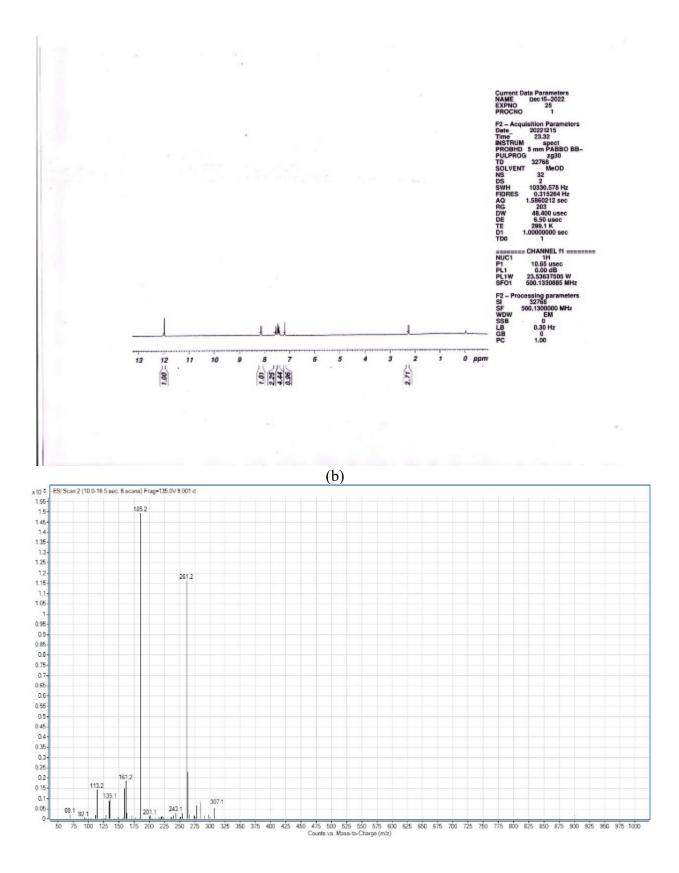
Figure S1. Chemical structure of 3-Phenyl-1-(Pyridine-4-Carbonyl)-1H-Pyrazole-4-Carboxylic Acid (PPP), 3-(4-Acetophenyl-)1-(Pyridine-4-Carbonyl)-1H-Pyrazole-4-Carboxylic Acid (APP) and 3-(4-Methylphenyl)-1-(Pyridine-4-Carbonyl)-1H-Pyrazole-4-Carboxylic Acid (MPP) derivatives

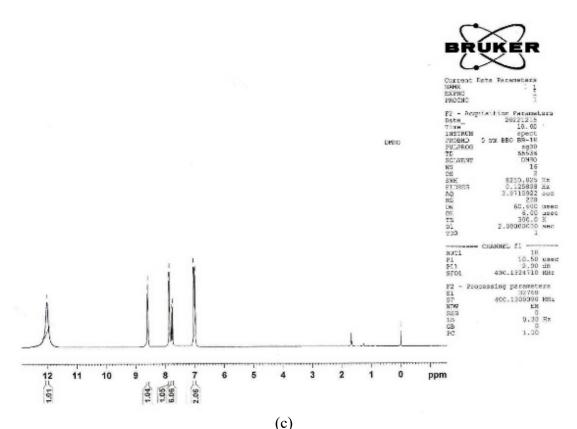




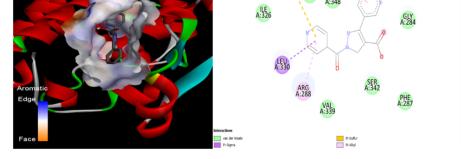








(c) Figure S2. NMR and mass spectrometry data for all derivatives (a) PPP, (b) APP, (c) MPP, obtained through experimental



PDB ID : 1IW0

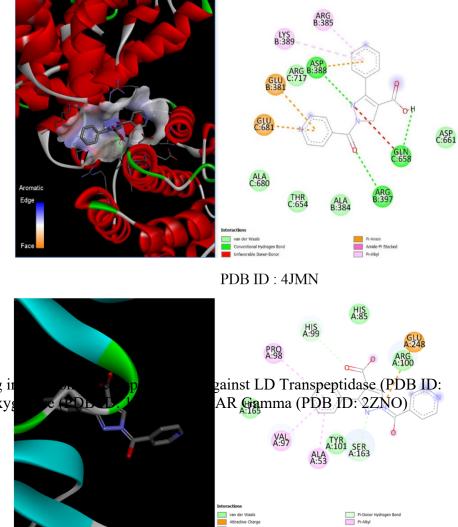
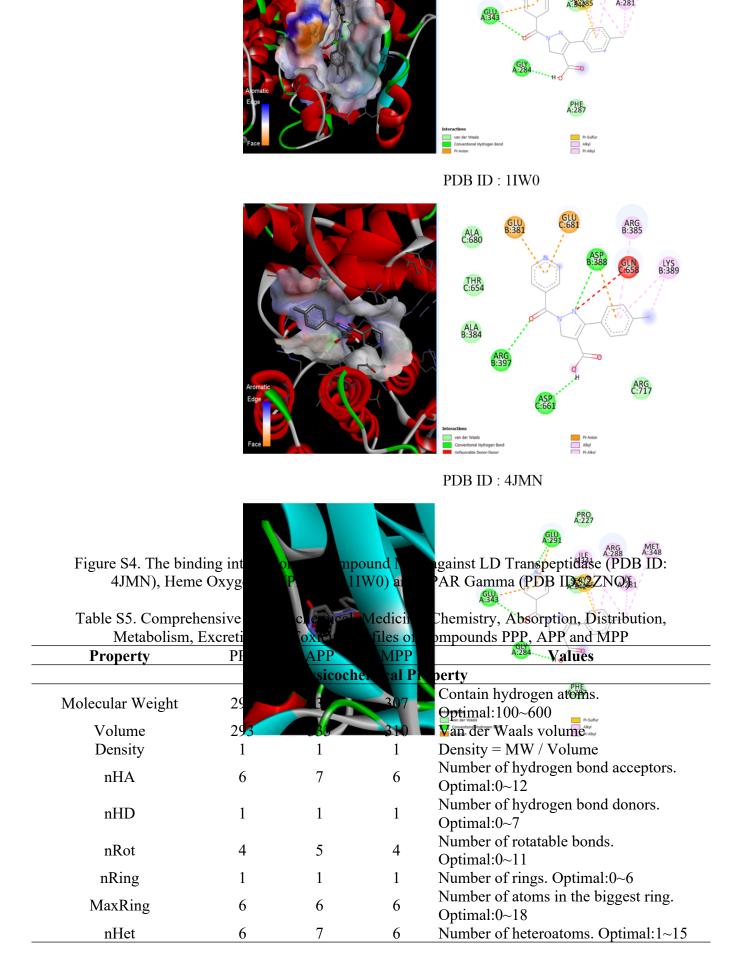


Figure S3. The binding in 4JMN), Heme Oxys



fChar	0.0	0.0	0.0	Formal charge. Optimal:-4~4
nRig	19	20	19	Number of rigid bonds. Optimal:0~30
Flexibility	0.0	0.30	0.20	Flexibility = nRot / nRig
Stereo Centers	0.0	0.0	0.0	Stereo Centers. Optimal: 2
TPSA	85.00	102.15	85.08	Topological Polar Surface Area.
				Optimal:0~140
logS	-3.00	-3.70	-4.41	The logarithm of aqueous solubility
2				value. The logarithm of the n-octanol/water
logP	1.27	1.32	1.98	distribution coefficients at pH=7.4.
				The logarithm of the n-octanol/water
logD	1.49	1.55	1.86	distribution coefficient.
				Acid-base dissociation constant (pKa)
pka (Acid)	3.55	4.09	4.04	value represents the strength of a drug
P ()				molecule's acidity or basicity.
				Acid-base dissociation constant (pKa)
pka (Base)	1.26	1.53	1.65	value represents the strength of a drug
				molecule's acidity or basicity.
				The predicted melting point of a
				compound is expressed in degrees
Melting point	230.94	250.84	247.79	Celsius (°C).
Menting point	250.71	230.04	211.19	Melting points below 25°C are classified
				as liquids, while melting points above
				25°C are classified as solids.
				The predicted melting point of a
	222.65	205.05	202 70	compound is expressed in degrees
Boiling point	323.65	305.95	293.70	Celsius (°C).
				A normal boiling point below 25°C is categorized as a gas.
		2 Medici	nal Chemis	0 0
		2. Miculti		A measure of drug-likeness based on
				the concept of desirability;
QED	0.80	0.74	0.80	• Attractive: $> 0.67;$
				■ unattractive: 0.49~0.67;
				■ too complex: < 0.34
				■ ES: Easy to synthesize; HS: Hard to
				synthesize;
GASA	0.0	0.0	0.0	■ The output value represents the
				probability of being difficult to
				synthesize, ranging from 0 to 1.
				 Synthetic accessibility score is
				designed to estimate ease of synthesis of
Synth	2.0	2.0	2.0	drug-like molecules.
J				■ SAscore 6, difficult to synthesize;
				SAscore <6,
				easy to synthesize

Fsp3	0.0	0.056	0.059	 The number of sp3 hybridized carbons / total carbon count, correlating with melting point and solubility. Fsp3 0.42 is considered a suitable value.
MCE-18	16.0	18.0	17.0	 MCE-18 stands for medicinal chemistry evolution. MCE-18 45 is considered a suitable value.
NPscore	-1.01	-0.91	-1.07	 Natural product-likeness score. This score is typically in the range from -5 to 5. The higher the score is, the higher the probability is that the molecule is a NP. MW 500; logP 5; Hacc 10; Hdon 5
Lipinski Rule	0.0	0.0	0.0	■ If two properties are out of range, a poor absorption or permeability is possible, one is acceptable.
Pfizer Rule	0.0	0.0	0.0	 logP > 3; TPSA < 75 Compounds with a high log P (>3) and low TPSA (<75) are likely to be toxic.
GSK Rule	0.0	0.0	0.0	 MW 400; logP 4 Compounds satisfying the GSK rule may have a more favorable ADMET profile
Golden Triangle	0.0	0.0	0.0	 200 MW 500; -2 logD 5 Compounds satisfying the Golden Triangle rule may have a more favorable ADMET profile.
PAINS	0 alerts	0 alerts	0 alerts	frequent hitters, Alpha-screen artifacts and reactive compound 480 substructures (J Med Chem 201052:2710,40)
ALARM NMR	0 alerts	0 alerts	0 alerts	201053:2719-40) Thiol reactive compounds.
BMS	0 alerts	0 alerts	0 alerts	undesirable, reactive compounds 176 substructures (J Chem Inf Model
Chelator Rule	0 alerts	0 alerts	0 alerts	200646:1060-8) Chelating compounds.
Colloidal aggregators	0.01	0.04	0.05	 Category 0: non-colloidal aggregators; Category 1: colloidal aggregators. The output value is the probability of being colloidal aggregators, within the range of 0 to 1.
FLuc inhibitors	0.17	0.27	0.34	■ Category 0: non-fLuc inhibitors;

Blue fluorescence	0.16	0.24	0.22	 Category 1: fLuc inhibitors. The output value is the probability of being fLuc inhibitors, within the range of 0 to 1. Category 0: non-blue fluorescence; Category 1: blue fluorescence. The output value is the probability of being blue fluorescence, within the range of 0 to 1. Category 0: non-green fluorescence; Category 1: green fluorescence.
Green fluorescence	0.25	0.35	0.35	 The output value is the probability of being green fluorescence, within the range of 0 to 1. Category 0: non-reactive compound; Category 1: reactive compound.
Reactive compounds	0.03	0.01	0.01	 The output value is the probability of being reactive compound, within the range of 0 to 1. Category 0: non-promiscuous compound; Category 1: promiscuous compound.
Promiscuous compounds	0.01	0.05	0.03	■ The output value is the probability of being promiscuous compound, within the range of 0 to 1.
Caco-2		3. Al	osorption	
Permeability	-5.12	-4.95	-4.60	Optimal: higher than -5.15 Log unit ■ low permeability: < 2 × 10-6 cm/s
MDCK Permeability	-4.60	-4.76	-4.70	 medium permeability: 2-20 × 10-6 cm/s high passive permeability: > 20 × 10-6 cm/s
PAMPA	0.98	0.98	0.99	 The experimental data for Peff was logarithmically transformed (logPeff). Molecules with log Peff values below 2.0 were classified as low-permeability (Category 0), while those with log Peff values exceeding 2.5 were classified as high-permeability (Category 1).
Pgp-inhibitor	0.02	0.02	0.04	 Category 1: Inhibitor; Category 0: Non-inhibitor; The output value is the probability of being

				Pgp-inhibitor ■ Category 1: substrate; ■ Category 0: Non-substrate;
Pgp-substrate	0.02	0.04	0.03	■ The output value is the probability of being
				Pgp-substrate Human Intestinal Absorption
				■ Category 1: HIA+(HIA < 30%);
HIA	0.00	0.01	0.00	■ Category 0: HIA-(HIA >= 30%);
				The output value is the probability of
				being HIA+
				 20% Bioavailability Category 1: F 20% + (bioavailability
				20%);
F20%	0.01	0.01	0.02	■ Category 0: F 20% - (bioavailability
12070	0.01	0.01	0.02	20%);
				■ The output value is the probability of being F 20%
				+
				■ 30% Bioavailability
				■ Category 1: F 30% + (bioavailability <
				30%);
F30%	0.05	0.03	0.02	■ Category 0: F 30% - (bioavailability 30%);
				■ The output value is the probability of
				being F 30%
				+
				 50% Bioavailability Catagory 1: E 50% + (bioavailability)
				■ Category 1: F 50% + (bioavailability < 50%);
	0.11	0.00		■ Category 0: F 50% - (bioavailability
F50%	() ()			
	0.11	0.08	0.06	50%);
	0.11	0.08	0.06	50%); ■ The output value is the probability of
	0.11	0.08	0.06	50%); ■ The output value is the probability of being F 50%
	0.11		0.06 tribution	50%); ■ The output value is the probability of
				 50%); The output value is the probability of being F 50% + Plasma Protein Binding
PPB	99.04			 50%); The output value is the probability of being F 50% + Plasma Protein Binding Optimal: < 90%.
PPB		4. Dis	tribution	 50%); The output value is the probability of being F 50% + Plasma Protein Binding Optimal: < 90%. Drugs with high protein-bound may
	99.04	4. Dis 95.68	tribution 98.71	 50%); The output value is the probability of being F 50% + Plasma Protein Binding Optimal: < 90%.
PPB VDss		4. Dis	tribution	 50%); The output value is the probability of being F 50% + Plasma Protein Binding Optimal: < 90%. Drugs with high protein-bound may have a low therapeutic index.
	99.04	4. Dis 95.68	tribution 98.71	 50%); The output value is the probability of being F 50% + Plasma Protein Binding Optimal: < 90%. Drugs with high protein-bound may have a low therapeutic index. Volume Distribution Optimal: 0.04-20L/kg Blood-Brain Barrier Penetration
	99.04	4. Dis 95.68	tribution 98.71	 50%); The output value is the probability of being F 50% + Plasma Protein Binding Optimal: < 90%. Drugs with high protein-bound may have a low therapeutic index. Volume Distribution Optimal: 0.04-20L/kg

Fu	0.80	3.31	1.01	 The fraction unbound in plasms Low: <5%; Middle: 5~20%; High: > 20%
				■ Category 0: Non-inhibitor; Category 1:
				inhibitor.
OATP1B1 inhibitor	0.64	0.81	0.84	The output value is the probability of
				being inhibitor, within the range of 0 to
				■ Category 0: Non-inhibitor; Category 1: inhibitor.
OATP1B3 inhibitor	0.71	0.92	0.68	■ The output value is the probability of
	0171	0.02	0.00	being inhibitor, within the range of 0 to
				1.
				■ Category 0: Non-inhibitor; Category 1:
BCRP				inhibitor.
inhibitor	0.04	0.02	0.01	The output value is the probability of
				being inhibitor, within the range of 0 to 1.
				■ Category 0: Non-inhibitor; Category 1:
				inhibitor.
MRP1 inhibitor	0.92	0.94	0.96	■ The output value is the probability of
				being inhibitor, within the range of 0 to
				1.
		5. Me	etabolism	
				 Category 1: Inhibitor; Category 0: Non-inhibitor;
CYP1A2 inhibitor	0.01	0.76	0.05	■ The output value is the probability of
				being inhibitor.
				■ Category 1: Substrate; Category 0:
CYP1A2 substrate	0.01	0.001	0.03	Non-substrate;
CTTTAZ Substrate	0.01	0.001	0.05	The output value is the probability of
				being substrate.
				 Category 1: Inhibitor; Category 0: Non-inhibitor;
CYP2C19 inhibitor	0.01	0.96	0.04	■ The output value is the probability of
				being inhibitor.
				■ Category 1: Substrate; Category 0:
CYP2C19 substrate	0.0	0.0	0.0	Non-substrate;
CIIZCI / Substrate	0.0	0.0	0.0	■ The output value is the probability of
	0.04	0.02	0.02	being substrate.
OVD0C0 1 1 1	0.84	0.92	0.93	■ Category 1: Inhibitor; Category 0:
CYP2C9 inhibitor				Non-inhibitor;
CYP2C9 inhibitor				
CYP2C9 inhibitor				■ The output value is the probability of
CYP2C9 inhibitor CYP2C9 substrate	0.07	0.0	0.06	

				■ The output value is the probability of being substrate.
CYP2D6 inhibitor	0.0	0.0	0.0	■ Category 1: Inhibitor; Category 0: Non-inhibitor;
				■ The output value is the probability of being inhibitor.
CYP2D6 substrate	0.0	0.0	0.0	■ Category 1: Substrate; Category 0: Non-substrate;
	0.01	0.0	0.07	■ The output value is the probability of being substrate.
CYP3A4 inhibitor	0.01	0.0	0.07	 Category 1: Inhibitor; Category 0: Non-inhibitor; The output value is the probability of
CYP3A4 substrate	0.0	0.0	0.0	 The output value is the probability of being inhibitor. Category 1: Substrate; Category 0:
C 11 SA4 substrate	0.0	0.0	0.0	 Category 1: Substrate, Category 0. Non-substrate; The output value is the probability of
CYP2B6 inhibitor	0.16	0.0	0.10	 The substrate is the presidently of being substrate. Category 1: Inhibitor; Category 0:
				Non-inhibitor; The output value is the probability of
CYP2B6 substrate	0.0	0.0	0.0	being inhibitor. ■ Category 1: Substrate; Category 0:
				Non-substrate; ■ The output value is the probability of
CYP2C8 inhibitor	0.80	1.0	1.0	being substrate. ■ Category 1: Inhibitor; Category 0:
				Non-inhibitor; ■ The output value is the probability of
				being inhibitor. ■ human liver microsomal (HLM)
				stability ■ Category 0: stable+ (HLM > 30 min); Category 1: unstable (HLM = 20 min);
HLM Stability	0.26	0.90	0.34	Category 1: unstable- (HLM 30 min). The output value is the probability of human liver microsomal instability,
				where a value closer to 1 indicates a higher likelihood of instability. The
				range is between 0 and 1.
		6. E	xcretion	
				■ The unit of predicted CLplasma
CLplasma	1.63	1.88	1.70	penetration is ml/min/kg. >15 ml/min/kg: high clearance; 5-15
CLpiasilia	1.03	1.00	1./0	ml/min/kg: moderate clearance; < 5
T1/2	1.42	1.32	1.31	ml/min/kg: low clearance. ■ The unit of predicted T1/2 is hours.

				■ ultra-short half-life drugs: 1/2 < 1 hour; short
				half-life drugs: T1/2 between 1-4 hours; intermediate short half-life drugs: T1/2
				between 4-8 hours; long half-life drugs:
				T1/2 > 8 hours.
		7. 1	Foxicity	11/2 - 0 110013.
			<u>onreitej</u>	■ Molecules with IC50 10 M or 50%
				inhibition at
				10 M were classified as hERG+
				(Category 1),
hERG Blockers	0.09	0.10	0.10	■ while molecules with IC50 >10 M or
				< 50% inhibition at 10 M were
				classified as hERG - (Category 0).
				■ The output value is the probability of
				being hERG+, within the range of 0 to
				■ Molecules with IC50 10 M are
				classified as hERG+ (Category 1),
LEDC Dississing (10mm)	0.10	0.07	0.10	■ and molecules with IC50 > 10 M are classified as
hERG Blockers (10um)	0.10	0.07	0.10	hERG- (Category 0).
				■ The output value is the probability of
				being hERG+, within the range of 0 to
				■ Drug Induced Liver Injury.
				■ Category 1: drugs with a high risk of
				DILI;
DILI	0.98	0.99	0.98	■ Category 0: drugs with no risk of
				DILI.
				■ The output value is the probability of
				being toxic.
				AMES Toxicity
		0.40	0.04	■ Category 1: Ames positive(+);
AMES Muta genicity	0.25	0.42	0.24	■ Category 0: Ames negative(-);
				■ The output value is the probability of
				being toxic. ■ Rat Oral Acute Toxicity.
				 Category 0: low-toxicity, > 500
				mg/kg;
Rat Oral Acute Toxicity	0.48	0.43	0.43	■ Category 1: high-toxicity; < 500
	0.10			mg/kg.
				■ The output value is the probability of
				being toxic, within the range of 0 to 1.
				■ FDA Maximum (Recommended)
FDAMDD	0.32	0.30	0.30	Daily Dose.
TDAMDD	0.32	0.30	0.30	■ Category 1: FDAMDD (+);
	0.02	0.00	0.50	 Category 1: FDAMDD (+); Category 0: FDAMDD (-);

				The output value is the probability of being positive.
				Category 1: Sensitizer;
Skin Sensiti zation	0.30	0.13	0.34	Category 0: Non-sensitizer.
				The output value is the probability of
				being toxic, within the range of 0 to 1.
				■ Category 1: carcinogens;
Carcinogeni city	0.60	0.75	0.62	Category 0: non-carcinogens;
8 9				The output value is the probability of
				being toxic.
				■ Eye Corrosion
Eye				■ Category 1: corrosives; Category 0:
Corrosion	0.0	0.0	0.0	noncorrosives;
Controlion				■ The output value is the probability of
				being corrosives.
				■ Eye Irritation
Eye				■ Category 1: irritants; Category 0:
Irritation	0.63	0.59	0.66	nonirritants;
initiation				■ The output value is the probability of
				being irritants.
				Category 1: respiratory toxicants;
Respiratory	0.71	0.78	0.70	Category 0: non-respiratory toxicants
Respiratory	0.71	0.78	0.70	■ The output value is the probability of
				being toxic, within the range of 0 to 1.
				Human Hepatotoxicity
				■ Category 1: H-HT positive(+);
Human Hep atotoxicity	0.71	0.74	0.71	■ Category 0: H-HT negative(-);
				■ The output value is the probability of
				being toxic.
				■ Category 0: non-nephrotoxic (-);
				■ Category 1: nephrotoxic (+).
Drug-induce d	0.77	0.82	0.79	■ The output value is the probability of
Nephrotox icity				being nephrotoxic (+), within the range
				of 0 to 1.
				■ Category 0: non-ototoxicity (-);
				■ Category 1: ototoxicity (+).
Ototoxicity	0.59	0.73	0.64	■ The output value is the probability of
5				being ototoxicity (+), within the range of
				0 to 1.
				■ Category 0: non-hematotoxicity (-);
				■ Category 1: hematotoxicity (+).
Hematotoxic ity	0.43	0.60	0.47	The output value is the probability of
	0.10	0.00	,	being hematotoxicity (+), within the
				range of 0 to 1.
		_	_	■ Category 0: non-Genotoxicity (-);
Genotoxicity	0.91	0.97	0.87	■ Category 1: Genotoxicity (+).

NR-AhR	0.63	0.54	0.72	 Aryl hydrocarbon receptor Category 1: actives ; Category 0: inactives;
LC50DM	3.90	3.89 9. Tox	3.94 21 pathwa	lethal concentration. ■ The unit is log10[(mg/L)/(1000*MW)]
LC50FM	3.36	3.27	3.36	 96-hour fathead minnow 50 percent lethal concentration. The unit is log10[(mg/L)/(1000*MW)] 48-hour daphnia magna 50 percent
IGC50	2.87	2.69	2.87	 log10[(mg/L)/(1000*MW)] ■ Tetrahymena pyriformis 50 percent growth inhibition concentration. ■ The unit is log10[(mg/L)/(1000*MW)]
Bioconcentration Factors	0.18	0.08	0.18	 Bioconcentration factors are used for considering secondary poisoning potential and assessing risks to human health via the food chain. The unit is
		8. Environ	<u>mental t</u> ox	
Drug-induce d Neurotoxi city	0.75	0.72	0.78	 Category 0: non-neurotoxic (-); Category 1: neurotoxic (+). The output value is the probability of being neurotoxic (+), within the range of 0 to 1.
Hek293 Cytotoxicity	0.06	0.07	0.05	 Category 0: non-cytotoxicity (-); Category 1: cytotoxicity (+). The output value is the probability of being ototoxicity (+), within the range of 0 to 1.
A549 Cytotoxicity	0.01	0.01	0.01	 0 to 1. Category 0: non-cytotoxicity (-); Category 1: cytotoxicity (+). The output value is the probability of being ototoxicity (+), within the range of 0 to 1.
RPMI-8226 Immunitoxici ty	0.03	0.07	0.03	 The output value is the probability of being ototoxicity (+), within the range of 0 to 1. Category 0: non-cytotoxicity (-); Category 1: cytotoxicity (+). The output value is the probability of being ototoxicity (+), within the range of 0 to 1.

				■ The output value is the probability of
				being active.
				 Androgen receptor
				■ Category 1: actives ;
NR-AR	0.09	0.09	0.11	■ Category 0: inactives;
				The output value is the probability of being active.
				 Androgen receptor ligand-binding domain
NR-AR-LBD	0.01	0.0	0.0	■ Category 1: actives ;
INK-AK-LDD	0.01	0.0	0.0	■ Category 0: inactives;
				■ The output value is the probability of
				being active.
				■ Category 1: actives ;
NR-Aromatase	0.16	0.02	0.04	■ Category 0: inactives;
INIX-ATOIIIatase	0.10	0.02	0.04	■ The output value is the probability of
				being active.
				Estrogen receptor
				■ Category 1: actives ;
NR-ER	0.24	0.16	0.25	■ Category 0: inactives;
				■ The output value is the probability of
				being active.
				 Estrogen receptor ligand-binding
				domain
NR-ER-LBD	0.0	0.0	0.00	■ Category 1: actives ;
MR-LR-LDD	0.0	0.0	0.00	■ Category 0: inactives;
				The output value is the probability of being active.
				Peroxisome proliferator-activated
				receptor gamma
NR-PPAR-gam ma	0.23	0.03	0.08	■ Category 1: actives ;
NR-11 AR-gain ina	0.23	0.05	0.08	■ Category 0: inactives;
				The output value is the probability of
				being active.
				Antioxidant response element
				■ Category 1: actives ;
SR-ARE	0.64	0.21	0.38	■ Category 0: inactives;
				The output value is the probability of
				being active.
				■ ATPase family AAA domain-
				containing protein 5
SR-ATAD5	0.05	0.03	0.05	■ Category 1: actives ;
				Category 0: inactives; The sector tended in the number of the sector tended.
				■ The output value is the probability of
CD LICE	0.15	0.02	0.11	being active.
SR-HSE	0.15	0.02	0.11	Heat shock factor response element

SR-MMP	0.49	0.08	0.12	 Category 0: inactives; The output value is the probability of being active. Mitochondrial membrane potential Category 1: actives; Category 0: inactives; The output value is the probability of being active. p53, a tumor suppressor protein
SR-p53	0.06	0.02	0.03	 Category 1: actives ; Category 0: inactives; The output value is the probability of being active.
		10. Toxico	ophore R	6
Acute Toxicity Rule		0		 20 substructures; acute toxicity during oral administration
Genotoxic Carcinogenicity Rule		0		117 substructures;carcinogenicity or mutagenicity
NonGenotoxic Carcinogenicity Rule		0		 23 substructures; carcinogenicity through nongenotoxic mechanisms
Skin Sensitization Rule		1 alerts		155 substructures;skin irritation
Aquatic Toxicity Rule		0		99 substructures;toxicity to liquid(water)
NonBiodegradable Rule		1 alerts		19 substructures;non-biodegradable
SureChEMBL Rule		0		164 substructures;MedChem unfriendly status
Toxicophores Rule		1 alerts		154 toxic substructures from FAF-Drug4