
Support Information for:

eMolTox: prediction of molecular toxicity with confidence

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Text S1. Conformal Prediction

Conformal prediction evaluates the similarity between the new samples and the training data. The output represents the probability that the new molecule is either active, inactive or uncertain (in the case of the query molecule being outside the applicability domain of the model), given a user-defined significance level that sets the maximum allowable fraction of erroneous predictions. For the classification task, the problem of conformal prediction is how to estimate the confidence of predicting the class label. This is done by finding how strange (nonconformal) a new example is in comparison to the training set, by calculating the nonconformity measure. This is done by the following several steps:

First, we divide the training set Z into two disjoint subsets: a proper training set Z_t and a calibration set Z_c .

Second, fit a model h using the training set Z_t . The underlying model is RandomForest.

Third, calculate the nonconformity score for each sample in the calibration set Z_c . To construct a conformal predictor's prediction, we need to define a nonconformity measure. The nonconformity function, we use is

$$\alpha(i) = 1 - P_h(y_i/x_i),$$

where the (y_i/x_i) is the predicted probability of a sample classified as a given class using the model h we constructed in the second step.

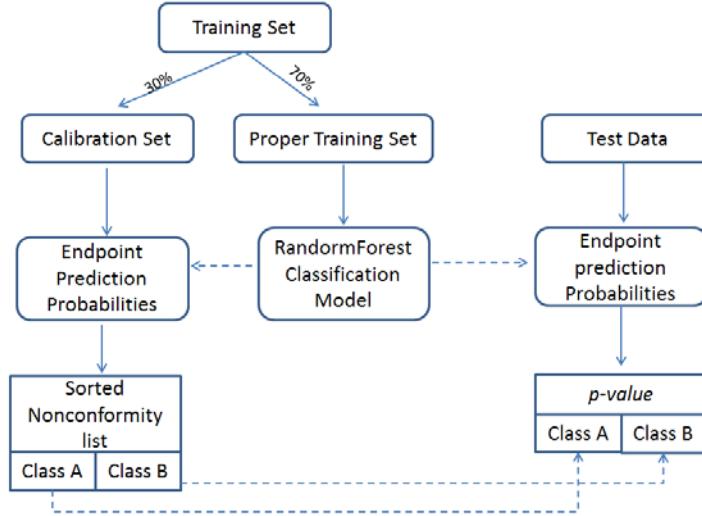
Fourth, to assess how different a new example x_n is from old examples, we need to compare nonconformity α_n to the nonconformity scores α_j of previous examples $x_j, j=1, \dots, k$ in the calibration sets. To make this comparison, we compute the fraction:

$$|\{j = 1, \dots, k: \alpha_j \geq \alpha_n\}|/n$$

i.e. the number of α_j that are as large or larger than α_n divided by the total number of α_j which gives the fraction of examples that have a larger α than the new sample under investigation. This fraction is called the *p-value* (not to be confused with the traditional p-values from statistics) for the new sample (x_n, y_n) . If the p-value is

small, then (x_n, y_n) is very nonconforming, i.e. the new example is different from the previous examples because of its high α score compared to most of the examples in the calibration set. On the other hand, if the p-value is large, then (x_n, y_n) is very conforming, i.e. very similar to the previous examples.

The following picture show a flow chart of conformal prediction:



For a given compound, a conformal predictor gives the p-value for active and inactive classes as p_1 and p_0 respectively. The output label under significance level \mathcal{E} (maximum allowable fraction of erroneous prediction) can be defined as:

Active: $p_1 > \mathcal{E}$ and $p_0 \leq \mathcal{E}$

Inactive: $p_0 > \mathcal{E}$ and $p_1 \leq \mathcal{E}$

Uncertain: $p_1 > \mathcal{E}$ and $p_0 > \mathcal{E}$, $p_1 \leq \mathcal{E}$ and $p_0 \leq \mathcal{E}$

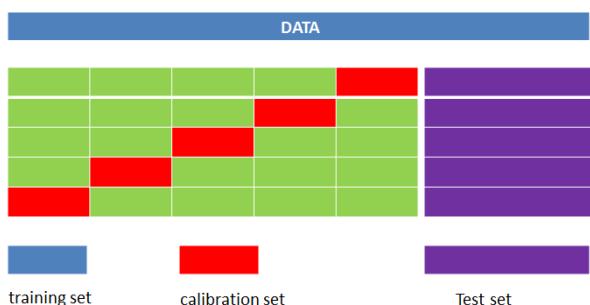
The associated output confidence for single labeled compound is defined as:

Confidence for compound labeled as Active: $1 - p_0$

Confidence for compound labeled as inactive: $1 - p_1$

Text S2. Model Construction and Performance Evaluation.

To evaluate the performance of each model, each data set was randomly divided into the training (70%) and test (30%) set. The training set was then further divided into a five-fold training set and calibrating set (see following figure). Each fold was used once as calibration set. The remaining training data were used to build underlying machine learning models. The averaged *p*-values estimated from five folds were used for decision making(Sun, et al., 2017). The Random Forest (RF) module of the Scikit-Learn package (Pedregosa, et al., 2011) was used as underlying machine learning models to build binary classification RF models. The Inductive Conformal Prediction module of nonconformist package (Linusson, 2017) was used for building Mondrial conformal predictor.



References:

- Linusson, H. (2017) Nonconformist, <https://github.com/donlnz/nonconformist>.
- Pedregosa, F., et al. (2011) Scikit-learn: Machine Learning in Python, *J. Mach. Learn. Res.*, **12**, 2825-2830.
- Sun, J., et al. (2017) Applying Mondrian Cross-Conformal Prediction To Estimate Prediction Confidence on Large Imbalanced Bioactivity Data Sets, *Journal of Chemical Information and Modeling*, **57**, 1591-1598.

Text S3. List of RDKit physicochemical descriptors used in this study.

'BalabanJ', 'BertzCT', 'Chi0', 'Chi0n', 'Chi0v', 'Chi1', 'Chi1n', 'Chi1v', 'Chi2n', 'Chi2v', 'Chi3n', 'Chi3v', 'Chi4n', 'Chi4v', 'EState_VSA1', 'EState_VSA10', 'EState_VSA11', 'EState_VSA2', 'EState_VSA3', 'EState_VSA4', 'EState_VSA5', 'EState_VSA6', 'EState_VSA7', 'EState_VSA8', 'EState_VSA9', 'ExactMolWt', 'FractionCSP3', 'HallKierAlpha', 'HeavyAtomCount', 'HeavyAtomMolWt', 'Ipc', 'Kappa1', 'Kappa2', 'Kappa3', 'LabuteASA', 'MaxAbsEStateIndex', 'MaxAbsPartialCharge', 'MaxEStateIndex', 'MaxPartialCharge', 'MinAbsEStateIndex', 'MinAbsPartialCharge', 'MinEStateIndex', 'MinPartialCharge', 'MolLogP', 'MolMR', 'MolWt', 'NHOHCount', 'NOCount', 'NumAliphaticCarbocycles', 'NumAliphaticHeterocycles', 'NumAliphaticRings', 'NumAromaticCarbocycles', 'NumAromaticHeterocycles', 'NumAromaticRings', 'NumHAcceptors', 'NumHDonors', 'NumHeteroatoms', 'NumRadicalElectrons', 'NumRotatableBonds', 'NumSaturatedCarbocycles', 'NumSaturatedHeterocycles', 'NumSaturatedRings', 'NumValenceElectrons', 'PEOE_VSA1', 'PEOE_VSA10', 'PEOE_VSA11', 'PEOE_VSA12', 'PEOE_VSA13', 'PEOE_VSA14', 'PEOE_VSA2', 'PEOE_VSA3', 'PEOE_VSA4', 'PEOE_VSA5', 'PEOE_VSA6', 'PEOE_VSA7', 'PEOE_VSA8', 'PEOE_VSA9', 'RingCount', 'SMR_VSA1', 'SMR_VSA10', 'SMR_VSA2', 'SMR_VSA3', 'SMR_VSA4', 'SMR_VSA5', 'SMR_VSA6', 'SMR_VSA7', 'SMR_VSA8', 'SMR_VSA9', 'SlogP_VSA1', 'SlogP_VSA10', 'SlogP_VSA11', 'SlogP_VSA12', 'SlogP_VSA2', 'SlogP_VSA3', 'SlogP_VSA4', 'SlogP_VSA5', 'SlogP_VSA6', 'SlogP_VSA7', 'SlogP_VSA8', 'SlogP_VSA9', 'TPSA', 'VSA_EState1', 'VSA_EState10', 'VSA_EState2', 'VSA_EState3', 'VSA_EState4', 'VSA_EState5', 'VSA_EState6', 'VSA_EState7', 'VSA_EState8', 'VSA_EState9', 'fr_Al_COO', 'fr_Al_OH', 'fr_Al_OH_noTert', 'fr_ArN', 'fr_Ar_COO', 'fr_Ar_N', 'fr_Ar_NH', 'fr_Ar_OH', 'fr_COO', 'fr_COO2', 'fr_C_O', 'fr_C_O_noCOO', 'fr_C_S', 'fr_HOCCN', 'fr_Imine', 'fr_NHO', 'fr_NH1', 'fr_NH2', 'fr_N_O', 'fr_Ndealkylation1', 'fr_Ndealkylation2', 'fr_Nhpyrrole', 'fr_SH', 'fr_aldehyde', 'fr_alkyl_carbamate', 'fr_alkyl_halide', 'fr_allylic_oxid', 'fr_amide', 'fr_amidine', 'fr_aniline', 'fr_aryl_methyl', 'fr_azide', 'fr_azo', 'fr_barbitur', 'fr_benzene', 'fr_benzodiazepine', 'fr_bicyclic', 'fr_diazo', 'fr_dihydropyridine', 'fr_epoxide', 'fr_ester', 'fr_ether', 'fr_furan', 'fr_guanido', 'fr_halogen', 'fr_hdrzine', 'fr_hdrzone', 'fr_imidazole', 'fr_imide', 'fr_isocyan', 'fr_isothiocyan', 'fr_ketone', 'fr_ketone_Topliss', 'fr_lactam', 'fr_lactone', 'fr_methoxy', 'fr_morpholine', 'fr_nitrile', 'fr_nitro', 'fr_nitro_arom', 'fr_nitro_arom_nonortho', 'fr_nitroso', 'fr_oxazole', 'fr_oxime', 'fr_para_hydroxylation', 'fr_phenol', 'fr_phenol_noOrthoHbond', 'fr_phos_acid', 'fr_phos_est', 'fr_piperdine', 'fr_piperazine', 'fr_priamide', 'fr_prisulfonam', 'fr_pyridine', 'fr_quatN', 'fr_sulfide', 'fr_sulfonam', 'fr_sulfone', 'fr_term_acetylene', 'fr_tetrazole', 'fr_thiazole', 'fr_thiocyan', 'fr_thiophene', 'fr_unbrch_alkane', 'fr_urea'

Table S1. Details of data sets for toxicity predictor construction. Data sets were collected from ChEMBL, Pubchem, Toxnet, eChemPortal databases and literature. For off-target data sets, compounds and their corresponding bioactivity information were derived from CHEMBL database and bioactivity values of 1 μ M or lower is classified as active. For those targets with very few inactive compounds, we borrow inactive compounds from their homology neighbors.

Model_ID	Data Description	Potential Injury	N_active	N_inact
M_001	Modulator of Sodium channel protein type V alpha subunit	Heart	122	610
M_002	Modulator of Calcitonin gene-related peptide type 1 receptor	Heart, Nervous system, Gastrointestinal, blood	554	2451
M_003	Modulator of Receptor protein-tirosine kinase erbB-2	Heart	1035	5171
M_004	Modulator of Dopamine D1 receptor	Central nervous system, Kidney, Heart	764	3824
M_005	Modulator of Sodium channel protein type IX alpha subunit	Heart	2159	8554
M_006	Modulator of P2X purinoceptor 3	Nervous system, immune, Kidney	848	565
M_007	Modulator of Endothelin receptor ET-B	Heart, respiratory, blood, Kidney, Gastrointestinal	513	1077
M_008	Modulator of Neuronal acetylcholine receptor protein alpha-4 subunit	Central nervous system	178	890
M_009	Modulator of Serotonin transporter	Central nervous system, blood, Heart, Gastrointestinal	3502	11228
M_010	Modulator of Adenosine A2a receptor	Heart, blood, Central nervous system	2997	12876
M_011	Modulator of Bradykinin B2 receptor	Heart, respiratory, Kidney, Central nervous system	383	1928
M_012	Modulator of Neuronal acetylcholine receptor; alpha3/beta4	Central nervous system	230	1150
M_013	Modulator of Serotonin 2c (5-HT2c) receptor	Nervous system	2325	11615
M_014	Modulator of HMG-CoA reductase	Kidney	192	959
M_015	Modulator of Cholecystokinin A receptor	Gastrointestinal, gallbladder, Nervous system	265	1325
M_016	Modulator of TNF-alpha	Immune	349	1745
M_017	Modulator of P2X purinoceptor 7	Nervous system, immune, Kidney	1872	4075
M_018	Modulator of Serotonin 2a (5-HT2a) receptor	Nervous system, blood, Heart	3086	13014
M_019	Modulator of Beta-1 adrenergic receptor	Heart	883	4410
M_020	Modulator of Platelet-derived growth factor receptor	Heart	171	854
M_021	Modulator of GABA-A receptor; anion channel	Central nervous system	974	2739
M_022	Modulator of Vascular endothelial growth factor receptor 1	Heart	1041	5198
M_023	Modulator of Platelet-derived growth factor receptor alpha	Heart	331	1655
M_024	Modulator of Melatonin receptor 1B	Central nervous system	739	1540
M_025	Modulator of Delta opioid receptor	Central nervous system	2398	11186
M_026	Modulator of GABA-A receptor; alpha-2/beta-3/gamma-2	Central nervous system	492	1888
M_027	Modulator of Peroxisome proliferator-activated receptor gamma	Kidney, Heart, immune	2099	10499
M_028	Modulator of Adenosine A1 receptor	Nervous system, Kidney, Heart	2584	12934
M_029	Modulator of Muscarinic acetylcholine receptor M4	Nervous system, Heart	448	2240
M_030	Modulator of Vascular endothelial growth factor receptor	Heart	141	705
M_031	Modulator of Glucocorticoid receptor	Endocrine, immune, Nervous system	1875	9373
M_032	Modulator of Cyclooxygenase-2	Kidney	1553	7794
M_033	Modulator of Kappa opioid receptor	Nervous system, Heart, Gastrointestinal	2831	13273
M_034	Modulator of Platelet activating factor receptor	Immune, respiratory, blood, Heart, Kidney	310	688

M_035	Modulator of Sigma opioid receptor	Nervous system	1862	9303
M_036	Modulator of Androgen Receptor	Endocrine, Central nervous system	1378	6890
M_037	Modulator of Neuropeptide Y receptor type 1	Gastrointestinal, immune, Nervous system	398	1990
M_038	Modulator of Monoamine oxidase A	Pharmacokinetics, Central nervous system	450	2250
M_039	Modulator of Dopamine D2 receptor	Central nervous system, Gastrointestinal, Heart, Kidney	4693	11918
M_040	Modulator of Vascular endothelial growth factor receptor 2	Heart	4642	9422
M_041	Modulator of Alpha-1a adrenergic receptor	Heart	1437	7176
M_042	Modulator of Glutamate receptor ionotropic, AMPA	Central nervous system	135	675
M_043	Modulator of Prostanoid EP2 receptor	Immune, urogenital	162	810
M_044	Modulator of Serotonin 3a (5-HT3a) receptor	Gastrointestinal, Heart, Central nervous system	365	1824
M_045	Modulator of Vasopressin V1a receptor	Heart, Kidney, Central nervous system	533	2665
M_046	Modulator of Cannabinoid CB1 receptor	Nervous system, Heart	2844	11626
M_047	Modulator of Serotonin 2b (5-HT2b) receptor	Heart, Nervous system, Gastrointestinal	1111	5550
M_048	Modulator of Cannabinoid CB2 receptor	Immune, Nervous system	3674	11038
M_049	Modulator of Adenosine A2b receptor	Immune, Peripheral nervous system	1434	5003
M_050	Modulator of Angiotensin II type 2 (AT-2) receptor	Kidney	288	1224
M_051	Modulator of Alpha-1b adrenergic receptor	Heart, Nervous system, Kidney	1084	5416
M_052	Modulator of Phosphodiesterase 3	Heart, blood	179	895
M_053	Modulator of Type-1 angiotensin II receptor	Kidney, Heart	626	2367
M_054	Modulator of HERG	Heart	1606	8024
M_055	Modulator of Neurokinin 2 receptor	Gastrointestinal, immune, gonad	728	3639
M_056	Modulator of Mu opioid receptor	Nervous system, respiratory, Gastrointestinal, Heart, pancreas, Kidney	2783	12209
M_057	Modulator of GABA-A receptor; alpha-5/beta-3/gamma-2	Central nervous system	604	1887
M_058	Modulator of Alpha-2a adrenergic receptor	Heart, Nervous system, Gastrointestinal, pancreas	620	3104
M_059	Modulator of Muscarinic acetylcholine receptor M5	Nervous system	325	1624
M_060	Modulator of Serotonin 4 (5-HT4) receptor	Gastrointestinal, Heart, Central nervous system	445	2225
M_061	Modulator of Beta-3 adrenergic receptor	Heart	1324	6614
M_062	Modulator of Serotonin 1a (5-HT1a) receptor	Nervous system, Heart, endocrine	3257	11716
M_063	Modulator of Neurokinin 1 receptor	Nervous system, immune, Gastrointestinal, gonad	2214	9498
M_064	Modulator of Angiotensin-converting enzyme	Heart, Kidney	376	1880
M_065	Modulator of Urotensin II receptor	Kidney, Heart	292	1458
M_066	Modulator of Muscarinic acetylcholine receptor M3	Gastrointestinal, Nervous system, respiratory, Heart, liver	1363	6815
M_067	Modulator of Platelet-derived growth factor receptor beta	Heart	720	3598
M_068	Modulator of Neuronal acetylcholine receptor; alpha4/beta2	Central nervous system	592	2960
M_069	Modulator of Dopamine transporter	Nervous system	1795	8974
M_070	Modulator of Serotonin 1b (5-HT1b) receptor	Nervous system, Heart	907	4535
M_071	Modulator of Glutamate NMDA receptor	Central nervous system	267	1335
M_072	Modulator of Histamine H1 receptor	Immune, Nervous system, Heart, Gastrointestinal	910	4548
M_073	Modulator of Neuronal acetylcholine receptor protein alpha-7 subunit	Central nervous system	384	1920
M_074	Modulator of Muscarinic acetylcholine receptor M2	Heart, respiratory, Nervous system	1145	5722

M_075	Modulator of Histamine H2 receptor	Gastrointestinal, immune, Heart	118	590
M_076	Modulator of Acetylcholinesterase	Central nervous system	1846	9245
M_077	Modulator of Endothelin receptor ET-A	Heart, Kidney, Nervous system, Gastrointestinal	1099	1007
M_078	Modulator of Muscarinic acetylcholine receptor M1	Nervous system, respiratory, Gastrointestinal, Heart, liver	1186	5935
M_079	Modulator of GABA-A receptor; alpha-1/beta-3/gamma-2	Central nervous system	573	1910
M_080	Modulator of Beta-2 adrenergic receptor	Heart, blood	1095	5469
M_081	Modulator of GABA-A receptor; alpha-3/beta-3/gamma-2	Central nervous system	617	1889
M_082	Modulator of Adenosine A3 receptor	Nervous system, respiratory	2711	8890
M_083	Modulator of Serotonin 7 (5-HT7) receptor	Nervous system, immune	1417	7081
M_084	Modulator of Norepinephrine transporter	Heart, Nervous system	2232	11150
M_085	Modulator of Alpha-2b adrenergic receptor	Heart, Nervous system	353	1769
M_086	Modulator of Vascular endothelial growth factor receptor 3	Heart	364	1820
M_087	Modulator of Cyclooxygenase-1	Kidney	296	1479
M_088	Differential cytotoxicity (isogenic chicken DT40 cell lines)	Genotoxicity	1939	5132
M_089	Antagonist of the glucocorticoid receptor (GR) signaling pathway	Endocrine, immune, Nervous system	373	6143
M_090	Differential cytotoxicity (isogenic chicken DT40 Rev3 mutant cell line)	Genotoxicity	2063	4980
M_091	Agonist of H2AX	DNA damage	411	6814
M_092	Agonist of the glucocorticoid receptor (GR) signaling pathway	Endocrine, immune, Nervous system	178	7158
M_093	Induce genotoxicity in human embryonic kidney cells	Genotoxicity	284	7444
M_094	Cytotoxicity in HEK293 cells - 40 hour	Kidney	702	6482
M_095	Activators of the human pregnane X receptor (PXR) signaling pathway	Liver	236	1749
M_096	Inhibitors of Hepatocyte nuclear factor 4 (HNF4) dimerization	Liver	1733	513
M_097	Antagonist of the androgen receptor (AR) signaling pathway	Endocrine, Central nervous system	532	6207
M_098	Disruptors of the mitochondrial membrane potential	Liver	992	5409
M_099	Agonist of the antioxidant response element (ARE) signaling pathway	Liver	967	5113
M_100	Cytotoxicity in HepG2 cells - 40 hour	Liver	854	6282
M_101	Antagonist of the estrogen receptor alpha (ER-alpha) signaling pathway	Endocrine	376	6361
M_102	Inhibitors and Substrates of Cytochrome P450 3A4	Liver	3435	7013
M_103	Inhibitors and Substrates of Cytochrome P450 2C9	Liver	1266	6902
M_104	Antagonist of the farnesoid-X-receptor (FXR) signaling pathway	Liver	315	6464
M_105	Activator the aryl hydrocarbon receptor (AhR) signaling pathway	Liver	873	6352
M_106	Antagonist of the constitutive androstane receptor (CAR) signaling pathway	Liver	460	6823
M_107	Activators of the heat shock response signaling pathway	Liver	412	6370
M_108	Inhibitors and Substrates of Cytochrome P450 2D6	Liver	1616	6295
M_109	Differential cytotoxicity against isogenic chicken DT40 cell lines with known DNA damage response pathways - Rad54/Ku70 mutant cell line	Genotoxicity	2070	5230
M_110	Agonist of the p53 signaling pathway	DNA damage	510	6922
M_111	Antagonist of the thyroid receptor (TR) signaling pathway	Endocrine, Heart	370	5602
M_112	Antagonist of the peroxisome proliferator-activated receptor gamma (PPARg) signaling pathway	Kidney, Heart, immune	375	5856
M_113	Cytotoxicity in HepG2 cells - 16 hour	Liver	487	6820
M_114	Antagonist of the retinoic acid receptor (RAR) signaling	Immune	576	5095

	pathway			
M_115	Cytotoxicity in HEK293 cells - 32 hour	Kidney	583	6610
M_116	Cytotoxicity in HepG2 cells - 8 hour	Liver	329	7058
M_117	Antagonist of the vitamin D receptor (VDR) signaling pathway	Liver	364	6035
M_118	Antagonist of the estrogen receptor alpha (ER-alpha) signaling pathway	Endocrine	391	6508
M_119	Agonist of the androgen receptor (AR) signaling pathway	Endocrine, Central nervous system	249	7126
M_120	Agonist of the AP-1 signaling pathway	Cancer	599	6406
M_121	Cytotoxicity in HepG2 cells - 32 hour	Liver	758	6383
M_122	Cytotoxicity in HepG2 cells - 0 hour	Liver	111	7419
M_123	Cytotoxicity in HEK293 cells - 24 hour	Kidney	477	6792
M_124	Antagonist of the retinoid-related orphan receptor gamma (ROR-gamma) signaling pathway	Immune	529	4999
M_125	Agonist of the estrogen receptor alpha (ER-alpha) signaling pathway	Endocrine	422	7256
M_126	Agonist of the thyroid stimulating hormone receptor (TSHR) signaling pathway	Endocrine, Heart	330	6822
M_127	Agonist of the constitutive androstane receptor (CAR) signaling pathway	Liver	893	6092
M_128	Agonist of the RXR signaling pathway	Liver	217	5865
M_129	Antagonist of the farnesoid-X-receptor (FXR) signaling pathway	Liver	226	6154
M_130	Cytotoxicity in HEK293 cells - 8 hour	Kidney	282	7062
M_131	Substrates of Cytochrome P450 2C19	Liver	1895	6400
M_132	Activators of Cytochrome P450 3A4	Liver	165	12631
M_133	Antagonist of the androgen receptor (AR) signaling pathway	Endocrine, Central nervous system	406	6256
M_134	Inhibit CYP1A2 Activity	Liver	4164	3639
M_135	Cytotoxicity in HEK293 cells - 16 hour	Heart	343	6982
M_136	Agonist of the peroxisome proliferator-activated receptor gamma (PPARg) signaling pathway	Kidney, Heart, immune	216	6851
M_137	Cytotoxicity in HepG2 cells - 24 hour	Liver	648	6547
M_138	Agonist of the farnesoid-X-receptor (FXR) signaling pathway	Liver	446	5829
M_139	Aromatase inhibitors	Endocrine	323	6075
M_140	Antagonist of the constitutive androstane receptor (CAR) signaling pathway	Liver	166	5239
M_141	Activators of Cytochrome P450 2A9	Liver	1368	13678
M_142	Activator Alzheimer's amyloid precursor	Central nervous system	1987	19870
M_143	Modulators of myocardial damage	Heart	2272	22716
M_144	Inhibit CYP2C9 Acitivity	Liver	18731	18731
M_145	Inhibit CYP2C19 Activity	Liver	20295	20295
M_146	Acute Oral Toxicity (rodent, LD50 < 5mg/kg)	Rodent	751	11072
M_147	Induce Phospholipidosis	Liver	442	789
M_148	Block OATP1B1 Transporter	Liver	1640	200
M_149	Agonist of Liver X receptor beta	Liver	560	3415
M_150	Mutagenicity	Mutagenicity	3296	2752
M_151	Acute Oral Toxicity (rodent, LD50 < 300mg/kg)	Rodent	6640	5063
M_152	Acute Oral Toxicity (rodent, LD50 < 50mg/kg)	Rodent	2664	9069
M_153	Agonist of Liver X receptor alpha	Liver	436	3415
M_154	Block OATP1B3 transporter	Liver	1726	130
M_155	Block Bile Salt Export Pump	Liver	361	309

M_156	Acute Oral Toxicity (Rat, LD50 < 300mg/kg)	Rat	2700	9227
M_157	Acute Oral Toxicity (Rat, LD50 < 5mg/kg)	Rat	237	11690
M_158	Acute Oral Toxicity (Rat, LD50 < 50mg/kg)	Rat	1002	10925
M_159	Carcinogenic Potency (Salmonella Mutagenicity)	carcinogenicity	392	426
M_160	Carcinogenic Potency (SingleCellCall)	carcinogenicity	780	675
M_161	Carcinogenic Potency (Rat)	carcinogenicity	571	597
M_162	Acute Oral Toxicity (Mouse, LD50 < 300mg/kg)	Mouse	4623	20233
M_163	Acute Oral Toxicity (Mouse, LD50 < 5mg/kg)	Mouse	149	24707
M_164	Acute Oral Toxicity (Mouse, LD50 < 50mg/kg)	Mouse	844	24012
M_165	Acute Oral Toxicity (Rabbit, LD50 < 300mg/kg)	Rabbit	227	695
M_166	Drug Induced Liver Injury	Liver	780	1042
M_167	Developmental Toxicity	develop	712	127
M_168	Endocrine Disruption	reproduction	648	430
M_169	Eye Corrosion	eye	883	1412
M_170	Eye Irritation	eye	3861	1346
M_171	Fish Tox (high risk)	fish	637	1387
M_172	Fish Tox (moderate risk)	fish	1158	866
M_173	Skin Sensitization	skin	343	329
M_174	Aquatic Tox(T pyriformis)	T pyriformis	1143	307

Table S2. Summary of Structural Alerts used in eMolTox analysis. Alerts were collected from ChEMBL, ToxAlerts and literatures.

ID	Alert Name	Number of Alerts	Reference
0	Chelating agents	55	1. Agrawal A, Johnson SL, Jacobsen JA, Miller MT, Chen LH, Pellecchia M, Cohen SM. Chelator fragment libraries for targeting metalloproteinases. <i>ChemMedChem</i> . 2010 Feb 1;5(2):195-9.
1	Acute Aquatic Toxicity	54	1. Hermens JL. Electrophiles and acute toxicity to fish. <i>Environmental Health Perspectives</i> . 1990;87:219-225. 2. Henk J.M. Verhaar, Cees J. van Leeuwen, Joop L.M. Hermens, Classifying environmental pollutants, <i>Chemosphere</i> , Volume 25, Issue 4, 1992, Pages 471-491.
2	Covalent Bind With Protein	108	1. Enoch SJ, Ellison CM, Schultz TW, Cronin MTD. (2011). A review of the electrophilic reaction chemistry involved in covalent protein binding relevant to toxicity. <i>Crit Rev Toxicol</i> 41:783-802.
3	Respiratory Sensitization	52	1. Enoch SJ et al, Development of Mechanism-Based Structural Alerts for Respiratory Sensitization Hazard Identification. <i>Chem. Res. Toxicol.</i> , 2012, 25 (11), pp 2490–2498
5	Mitochondrial Toxicity	17	1. Nelms MD, Mellor CL, Cronin MT, Madden JC, Enoch SJ. Development of an <i>in silico</i> profiler for mitochondrial toxicity. <i>Chemical research in toxicology</i> . 2015 Sep 24;28(10):1891-902.
6	Non genotoxic carcinogenicity	22	1. Benigni R, Bossa C, Tcheremenskaia O. Nongenotoxic carcinogenicity of chemicals: mechanisms of action and early recognition through a new set of structural alerts. <i>Chemical reviews</i> . 2013 Mar 8;113(5):2940-57.
8	Skin sensitization	135	1. Payne MP, Walsh PT. Structure-activity relationships for skin sensitization potential: development of structural alerts for use in knowledge-based toxicity prediction systems. <i>Journal of chemical information and computer sciences</i> . 1994 Jan;34(1):154-61. 2. Barratt MD, Baskett DA, Chamberlain M, Admans GD, Langowski JJ. An expert system rulebase for identifying contact allergens. <i>Toxicology in vitro</i> . 1994 Oct 1;8(5):1053-60. 3. Enoch SJ, Madden JC, Cronin MT. Identification of mechanisms of toxic action for skin sensitisation using a SMARTS pattern based approach. <i>SAR and QSAR in Environmental Research</i> . 2008 Jul 1;19(5-6):555-78. 4. Gerner I, Barratt MD, Zinke S, Schlegel K, Schlede E. Development and prevalidation of a list of structure-activity relationship rules to be used in expert systems for prediction of the skin-sensitising properties of chemicals. <i>Alternatives to laboratory animals: ATLA</i> . 2004 Nov;32(5):487-509.
9	Alert from Top200 Drug	82	1. Stepan AF, Walker DP, Bauman J, Price DA, Baillie TA, Kalgutkar AS, Aleo MD. Structural alert/reactive metabolite concept as applied in medicinal chemistry to mitigate the risk of idiosyncratic drug toxicity: a perspective based on the critical examination of trends in the top 200 drugs marketed in the United States. <i>Chemical research in toxicology</i> . 2011 Jul 11;24(9):1345-410.
10	Genotoxic carcinogenicity mutagenicity	81	1. Ashby J, Tennant RW. Chemical structure, <i>Salmonella</i> mutagenicity and extent of carcinogenicity as indicators of genotoxic carcinogenesis among 222 chemicals tested in rodents by the US NCI/NTP. <i>Mutation Research/Genetic Toxicology</i> . 1988 Jan 1;204(1):17-15. 2. Kazius J, McGuire R, Bursi R. Derivation and validation of toxicophores for mutagenicity prediction. <i>Journal of medicinal chemistry</i> . 2005 Jan 13;48(1):312-20. 3. Bailey AB, Chanderbhan R, Collazo-Braier N, Cheeseman MA, Twaroski ML. The use of structure-activity relationship analysis in the food contact notification program. <i>Regulatory Toxicology and Pharmacology</i> . 2005 Jul 31;42(2):225-35. 4. Benigni R, Bossa C. Structure alerts for carcinogenicity, and the <i>Salmonella</i> assay system: a novel insight through the chemical relational databases technology. <i>Mutation Research/Reviews in Mutation Research</i> . 2008 Oct 31;659(3):248-61.
11	Kidney Toxicity	6	1. Pizzo F, Gadaleta D, Lombardo A, Nicolotti O, Benfenati E. Identification of structural alerts for liver and kidney toxicity using repeated dose toxicity data. <i>Chemistry Central Journal</i> . 2015 Dec 1;9(1):62.
12	Hepatotoxicity	36	1. Liu R, Yu X, Wallqvist A. Data-driven identification of structural alerts for mitigating the risk of drug-induced human liver injuries. <i>Journal of cheminformatics</i> . 2015 Dec 1;7(1):4. 2. Pizzo F, Gadaleta D, Lombardo A, Nicolotti O, Benfenati E. Identification of structural alerts for liver and kidney toxicity using repeated dose toxicity data. <i>Chemistry Central Journal</i> . 2015 Dec 1;9(1):62. 3. Hewitt M, Enoch SJ, Madden JC, Przybylak KR, Cronin MT. Hepatotoxicity: a scheme for generating chemical categories for read-across, structural alerts and insights into mechanism (s) of action. <i>Critical reviews in toxicology</i> . 2013 Aug 1;43(7):537-58.
13	Developmental and mitochondrial toxicity	12	1. Structural alerts for developmental toxicity and mitochondrial toxicity molecular initiating events (Lhasa), Mukesh,P et al. 2014

			1. Benigni R and Bossa C. Structure alerts for carcinogenicity, and the Salmonella assay system: a novel insight through the chemical relational databases technology. <i>Mutation Research</i> 2008, 659, 248-261. 2. Enoch SJ, Cronin MTD. A review of the electrophilic reaction chemistry involved in covalent DNA binding. <i>Crit Rev Toxicol</i> 2010;40(8):728-748. 3. Kalgutkar AS et al. A comprehensive listing of bioactivation pathways of organic functional groups. <i>Current Drug Metabolism</i> 2005; 6, 161-225. 4. Kazius J et al. Derivation and Validation of Toxicophores for Mutagenicity Prediction. <i>J Med Chem</i> 2005; 48, 312-320. 5. Mekenyen O et al. Identification of the Structural Requirements for Mutagenicity by Incorporating Molecular Flexibility and Metabolic Activation of Chemicals I: TA100 Model. <i>Chem Res Toxicol</i> 2004, 17, 753-766. 6. Mekenyen O et al. Identifying the Structural Requirements for Chromosomal Aberration by Incorporating Molecular Flexibility and Metabolic Activation of Chemicals. <i>Chem Res Toxicol</i> 2007, 20, 1927-1941. 7. Serafimova R et al. Identification of the Structural Requirements for Mutagenicity, by Incorporating Molecular Flexibility and Metabolic Activation of Chemicals. II. General Ames Mutagenicity Model. <i>Chem Res Toxicol</i> 2007, 20, 662-676.
14	Covalent Bind With DNA	111	1. Kalgutkar AS, Soglia JR. Minimising the potential for metabolic activation in drug discovery. <i>Expert opinion on drug metabolism & toxicology</i> . 2005 Jun 1;1(1):91-142.
15	Idiosyncratic toxicity Metabolic activation	32	1. Enoch SJ, Ellison CM, Schultz TW, Cronin MT. A review of the electrophilic reaction chemistry involved in covalent protein binding relevant to toxicity. <i>Critical reviews in toxicology</i> . 2011 Oct 1;41(9):783-802.
16	Potential electrophilic agents	119	

Table S3. Performance of Mondral Conformal Predictors for all 174 Data Sets at three significance levels.

Model_ID	$\varepsilon = 0.05$			$\varepsilon = 0.10$			$\varepsilon = 0.20$		
	efficiency	validity (positive set)	validity (negative set)	efficiency	validity (positive set)	validity (negative set)	efficiency	validity (positive set)	validity (negative set)
M_001	0.44	1.00	0.97	0.94	0.92	0.90	0.80	0.62	0.83
M_002	0.98	0.98	0.98	0.94	0.96	0.93	0.85	0.85	0.86
M_003	0.98	0.97	0.94	0.91	0.89	0.89	0.81	0.80	0.80
M_004	0.99	0.90	0.96	0.92	0.86	0.91	0.82	0.80	0.82
M_005	0.97	0.95	0.96	0.93	0.92	0.91	0.82	0.82	0.81
M_006	0.97	0.97	0.97	0.92	0.92	0.90	0.81	0.84	0.78
M_007	0.97	0.96	0.94	0.92	0.90	0.88	0.81	0.84	0.79
M_008	0.94	0.96	0.96	0.99	0.89	0.96	0.91	0.78	0.90
M_009	1.00	0.95	0.96	0.93	0.90	0.91	0.81	0.81	0.81
M_010	0.99	0.94	0.96	0.93	0.91	0.91	0.83	0.82	0.82
M_011	0.96	0.97	0.96	0.92	0.94	0.92	0.84	0.79	0.85
M_012	0.80	0.94	0.97	0.99	0.87	0.93	0.84	0.80	0.82
M_013	0.99	0.96	0.96	0.93	0.91	0.91	0.83	0.82	0.82
M_014	0.97	0.88	0.96	0.92	0.88	0.90	0.80	0.79	0.79
M_015	0.99	0.96	0.96	0.95	0.91	0.94	0.82	0.84	0.81
M_016	0.91	0.98	0.98	0.94	0.92	0.94	0.87	0.76	0.88
M_017	0.97	0.95	0.96	0.92	0.91	0.92	0.82	0.82	0.81
M_018	0.98	0.97	0.96	0.93	0.92	0.91	0.82	0.82	0.81
M_019	0.98	0.95	0.95	0.92	0.89	0.91	0.83	0.82	0.83
M_020	0.61	1.00	0.94	0.98	0.96	0.89	0.86	0.82	0.84
M_021	1.00	0.97	0.95	0.93	0.93	0.90	0.82	0.84	0.79
M_022	0.63	0.96	0.96	0.85	0.92	0.92	0.92	0.83	0.82
M_023	0.53	0.96	0.95	0.71	0.91	0.90	0.96	0.77	0.78
M_024	0.97	0.98	0.95	0.92	0.92	0.91	0.81	0.83	0.80
M_025	0.98	0.96	0.95	0.93	0.92	0.91	0.82	0.82	0.81
M_026	0.98	0.95	0.97	0.90	0.88	0.91	0.78	0.77	0.78
M_027	0.98	0.96	0.95	0.92	0.91	0.91	0.82	0.82	0.80
M_028	0.99	0.96	0.96	0.93	0.91	0.91	0.82	0.79	0.82
M_029	0.69	0.97	0.97	0.96	0.92	0.93	0.84	0.78	0.81
M_030	0.79	0.98	0.97	0.95	0.95	0.90	0.81	0.83	0.79
M_031	0.96	0.95	0.96	0.91	0.89	0.91	0.81	0.81	0.81
M_032	0.90	0.95	0.94	0.98	0.89	0.89	0.85	0.78	0.81
M_033	0.99	0.95	0.95	0.92	0.91	0.91	0.82	0.81	0.81
M_034	0.94	1.00	0.94	0.93	0.96	0.89	0.83	0.86	0.82
M_035	0.96	0.96	0.95	0.95	0.91	0.91	0.84	0.82	0.82
M_036	0.98	0.95	0.95	0.92	0.90	0.91	0.83	0.81	0.83
M_037	0.98	0.97	0.97	0.92	0.87	0.92	0.82	0.79	0.82
M_038	0.75	0.99	0.97	0.91	0.94	0.94	0.87	0.87	0.82

M_039	0.99	0.96	0.95	0.93	0.92	0.90	0.82	0.82	0.80
M_040	0.93	0.96	0.95	0.97	0.92	0.91	0.84	0.81	0.82
M_041	0.99	0.96	0.95	0.93	0.91	0.91	0.83	0.80	0.82
M_042	0.64	1.00	0.96	0.97	1.00	0.89	0.82	0.83	0.80
M_043	0.90	0.90	0.95	0.96	0.86	0.91	0.79	0.65	0.79
M_044	0.81	0.99	0.96	0.94	0.93	0.90	0.79	0.76	0.79
M_045	0.98	0.96	0.97	0.94	0.94	0.93	0.85	0.87	0.83
M_046	1.00	0.95	0.96	0.93	0.91	0.90	0.83	0.83	0.82
M_047	0.83	0.98	0.96	0.99	0.93	0.91	0.82	0.84	0.79
M_048	1.00	0.96	0.95	0.93	0.91	0.91	0.83	0.83	0.82
M_049	0.99	0.97	0.96	0.94	0.93	0.91	0.83	0.84	0.81
M_050	0.97	0.94	0.97	0.93	0.91	0.91	0.78	0.73	0.79
M_051	0.99	0.97	0.96	0.92	0.92	0.91	0.83	0.83	0.82
M_052	0.75	1.00	0.98	0.96	0.93	0.95	0.85	0.82	0.85
M_053	0.98	0.98	0.96	0.91	0.90	0.91	0.80	0.81	0.80
M_054	0.62	0.97	0.95	0.83	0.95	0.90	0.91	0.83	0.81
M_055	0.99	0.99	0.97	0.92	0.94	0.92	0.82	0.78	0.82
M_056	1.00	0.97	0.95	0.93	0.91	0.91	0.82	0.79	0.82
M_057	0.97	0.97	0.96	0.92	0.95	0.91	0.83	0.83	0.82
M_058	0.83	0.95	0.95	0.98	0.90	0.91	0.85	0.81	0.82
M_059	0.58	1.00	0.96	0.83	0.98	0.93	0.89	0.87	0.83
M_060	0.97	0.96	0.96	0.92	0.93	0.92	0.83	0.82	0.83
M_061	0.97	0.97	0.96	0.92	0.95	0.91	0.84	0.86	0.84
M_062	1.00	0.96	0.94	0.92	0.93	0.89	0.81	0.82	0.80
M_063	0.96	0.93	0.96	0.92	0.89	0.92	0.82	0.81	0.82
M_064	0.99	0.94	0.95	0.91	0.88	0.90	0.81	0.77	0.81
M_065	0.95	0.97	0.95	0.89	0.90	0.89	0.79	0.84	0.79
M_066	0.99	0.96	0.96	0.93	0.91	0.92	0.83	0.81	0.83
M_067	0.70	0.98	0.97	0.93	0.93	0.92	0.85	0.76	0.82
M_068	0.95	0.96	0.95	0.95	0.93	0.90	0.83	0.85	0.80
M_069	0.99	0.95	0.95	0.93	0.89	0.90	0.81	0.77	0.81
M_070	0.99	0.95	0.96	0.91	0.88	0.91	0.83	0.81	0.83
M_071	0.92	0.98	0.98	0.97	0.88	0.94	0.86	0.86	0.85
M_072	0.95	0.96	0.96	0.95	0.92	0.91	0.84	0.81	0.83
M_073	0.82	0.97	0.97	0.98	0.93	0.90	0.83	0.80	0.81
M_074	0.90	0.97	0.95	0.95	0.91	0.90	0.82	0.79	0.81
M_075	0.60	0.97	0.94	0.87	0.89	0.90	0.92	0.77	0.84
M_076	0.95	0.94	0.96	0.96	0.91	0.91	0.85	0.83	0.83
M_077	0.86	0.97	0.98	0.99	0.92	0.92	0.85	0.84	0.82
M_078	0.89	0.96	0.96	0.97	0.90	0.91	0.84	0.81	0.81
M_079	0.97	0.97	0.94	0.91	0.93	0.90	0.82	0.86	0.80
M_080	0.98	0.95	0.95	0.93	0.89	0.90	0.82	0.79	0.82
M_081	0.97	0.95	0.96	0.91	0.88	0.91	0.81	0.76	0.82
M_082	0.99	0.97	0.95	0.94	0.92	0.90	0.81	0.78	0.81

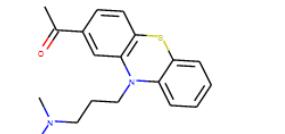
M_083	0.97	0.97	0.96	0.93	0.91	0.91	0.82	0.81	0.81
M_084	0.99	0.96	0.95	0.92	0.91	0.90	0.82	0.79	0.82
M_085	0.77	0.96	0.95	0.93	0.91	0.92	0.88	0.86	0.80
M_086	0.48	0.93	0.96	0.73	0.88	0.90	1.00	0.78	0.78
M_087	0.45	0.99	0.95	0.80	0.96	0.89	0.95	0.81	0.80
M_088	0.49	0.95	0.97	0.71	0.90	0.92	0.97	0.79	0.81
M_089	0.39	0.98	0.98	0.67	0.96	0.93	0.97	0.85	0.83
M_090	0.39	0.97	0.96	0.61	0.93	0.91	0.90	0.83	0.81
M_091	0.34	0.98	0.96	0.59	0.91	0.91	0.89	0.83	0.81
M_092	0.20	1.00	0.97	0.50	0.98	0.94	0.84	0.93	0.84
M_093	0.27	0.99	0.97	0.50	0.98	0.93	0.85	0.86	0.84
M_094	0.24	0.99	0.96	0.49	0.95	0.91	0.82	0.86	0.83
M_095	0.48	0.96	0.97	0.76	0.92	0.91	1.00	0.83	0.81
M_096	0.38	0.96	0.98	0.61	0.93	0.90	0.90	0.82	0.82
M_097	0.52	0.96	0.96	0.72	0.93	0.91	0.99	0.81	0.82
M_098	0.61	0.98	0.96	0.82	0.95	0.91	0.94	0.87	0.79
M_099	0.34	0.98	0.96	0.58	0.91	0.91	0.90	0.79	0.80
M_100	0.26	0.96	0.96	0.48	0.93	0.92	0.81	0.84	0.81
M_101	0.36	0.97	0.96	0.66	0.91	0.92	0.96	0.80	0.82
M_102	0.68	0.96	0.95	0.87	0.92	0.90	0.93	0.83	0.81
M_103	0.36	0.97	0.96	0.65	0.92	0.91	0.96	0.83	0.80
M_104	0.46	0.96	0.97	0.69	0.90	0.93	0.97	0.80	0.83
M_105	0.57	0.95	0.96	0.78	0.90	0.90	0.99	0.83	0.80
M_106	0.28	0.94	0.97	0.51	0.88	0.92	0.80	0.78	0.83
M_107	0.13	0.95	0.96	0.32	0.94	0.92	0.72	0.76	0.82
M_108	0.57	0.98	0.96	0.77	0.93	0.91	0.98	0.82	0.81
M_109	0.45	0.96	0.95	0.66	0.90	0.90	0.95	0.80	0.79
M_110	0.41	0.95	0.97	0.62	0.92	0.93	0.92	0.78	0.81
M_111	0.45	0.96	0.96	0.65	0.93	0.90	0.91	0.85	0.80
M_112	0.40	0.96	0.97	0.61	0.92	0.92	0.91	0.83	0.81
M_113	0.23	0.95	0.96	0.48	0.88	0.92	0.83	0.80	0.82
M_114	0.39	0.94	0.96	0.61	0.91	0.91	0.93	0.81	0.79
M_115	0.27	0.96	0.96	0.55	0.90	0.92	0.89	0.77	0.82
M_116	0.14	0.99	0.97	0.33	0.93	0.92	0.69	0.80	0.84
M_117	0.44	0.93	0.96	0.68	0.87	0.91	0.98	0.77	0.80
M_118	0.51	0.97	0.97	0.71	0.94	0.92	0.95	0.87	0.81
M_119	0.10	1.00	0.97	0.24	0.99	0.95	0.72	0.91	0.88
M_120	0.31	0.99	0.95	0.53	0.95	0.91	0.85	0.88	0.82
M_121	0.27	0.95	0.95	0.50	0.93	0.90	0.86	0.81	0.81
M_122	0.02	1.00	0.99	0.04	1.00	0.97	0.17	1.00	0.91
M_123	0.18	1.00	0.97	0.45	0.94	0.92	0.84	0.80	0.83
M_124	0.48	0.95	0.96	0.71	0.91	0.91	0.98	0.85	0.80
M_125	0.26	0.93	0.97	0.45	0.89	0.92	0.84	0.76	0.83
M_126	0.30	0.98	0.97	0.55	0.94	0.93	0.87	0.83	0.83

M_127	0.63	0.96	0.96	0.81	0.91	0.92	0.96	0.80	0.81
M_128	0.04	1.00	0.98	0.13	0.95	0.95	0.39	0.75	0.87
M_129	0.45	0.99	0.96	0.69	0.93	0.91	0.95	0.82	0.82
M_130	0.18	0.95	0.97	0.36	0.91	0.92	0.70	0.81	0.83
M_131	0.44	0.96	0.96	0.67	0.92	0.91	0.95	0.85	0.81
M_132	0.21	1.00	0.96	0.64	0.96	0.92	0.94	0.84	0.85
M_133	0.44	0.95	0.97	0.66	0.93	0.92	0.94	0.76	0.81
M_134	0.57	0.95	0.96	0.75	0.91	0.93	0.99	0.82	0.81
M_135	0.20	0.95	0.97	0.44	0.92	0.93	0.79	0.83	0.84
M_136	0.24	0.99	0.97	0.52	0.92	0.92	0.88	0.75	0.83
M_137	0.23	0.96	0.96	0.46	0.91	0.93	0.81	0.83	0.83
M_138	0.49	0.94	0.96	0.71	0.89	0.93	0.97	0.78	0.84
M_139	0.29	1.00	0.97	0.57	0.97	0.91	0.89	0.91	0.80
M_140	0.14	1.00	0.97	0.40	0.96	0.93	0.75	0.88	0.84
M_141	0.42	0.96	0.96	0.67	0.89	0.92	0.95	0.79	0.82
M_142	0.67	0.94	0.96	0.86	0.90	0.91	0.94	0.81	0.82
M_143	0.60	0.97	0.96	0.80	0.92	0.91	0.96	0.83	0.81
M_144	0.32	0.95	0.96	0.50	0.91	0.91	0.79	0.82	0.80
M_145	0.29	0.96	0.96	0.46	0.92	0.92	0.74	0.83	0.82
M_146	0.61	0.95	0.95	0.84	0.89	0.91	0.94	0.82	0.82
M_147	0.36	0.99	0.97	0.60	0.91	0.95	0.94	0.85	0.87
M_148	0.32	0.95	0.97	0.59	0.89	0.92	0.85	0.82	0.87
M_149	0.96	0.96	0.95	0.91	0.89	0.91	0.82	0.77	0.83
M_150	0.51	0.97	0.96	0.73	0.91	0.90	0.98	0.80	0.81
M_151	0.44	0.96	0.96	0.65	0.92	0.91	0.92	0.81	0.79
M_152	0.47	0.96	0.97	0.68	0.91	0.92	0.95	0.81	0.81
M_153	0.96	0.98	0.95	0.91	0.95	0.90	0.81	0.82	0.81
M_154	0.46	0.95	0.95	0.64	0.91	0.95	0.95	0.79	0.92
M_155	0.61	0.96	0.97	0.86	0.93	0.88	0.96	0.86	0.80
M_156	0.45	0.97	0.95	0.65	0.93	0.91	0.94	0.82	0.81
M_157	0.63	0.90	0.96	0.87	0.83	0.92	0.93	0.73	0.84
M_158	0.45	0.97	0.96	0.73	0.96	0.91	0.98	0.86	0.83
M_159	0.35	1.00	0.92	0.55	0.98	0.88	0.85	0.89	0.74
M_160	0.21	0.98	0.97	0.35	0.92	0.94	0.63	0.83	0.85
M_161	0.21	0.94	0.95	0.35	0.90	0.93	0.57	0.84	0.84
M_162	0.50	0.96	0.96	0.71	0.91	0.91	0.97	0.80	0.82
M_163	0.04	1.00	0.98	0.26	0.96	0.95	0.86	0.78	0.89
M_164	0.47	0.97	0.96	0.78	0.92	0.92	0.96	0.83	0.83
M_165	0.13	0.99	0.97	0.27	0.94	0.95	0.58	0.84	0.88
M_166	0.18	0.97	0.98	0.40	0.94	0.94	0.69	0.87	0.86
M_167	0.06	0.98	0.97	0.18	0.94	0.95	0.38	0.87	0.92
M_168	0.14	1.00	0.94	0.30	0.95	0.90	0.60	0.81	0.80
M_169	0.97	0.98	0.96	0.97	0.96	0.92	0.86	0.86	0.84
M_170	0.90	0.95	0.94	0.99	0.91	0.89	0.83	0.81	0.78

M_171	0.44	0.98	0.96	0.70	0.90	0.89	0.97	0.83	0.81
M_172	0.55	0.97	0.95	0.75	0.91	0.89	0.99	0.84	0.77
M_173	0.40	0.92	0.98	0.67	0.86	0.92	0.97	0.75	0.85
M_174	0.81	0.97	1.00	0.93	0.92	0.94	0.88	0.81	0.83

Figure S1. Example figure of result table from conformal prediction.

Molecule Depiction



Molecule properties:

Property	Value
Num Aliphatic Carbocycles	0
Num Heteroatoms	4
Num Hydrogen Bond Acceptor	4
Num Aromatic Carbocycles	2
LogP	4.444
Num Rotatable Bonds	5
Num Aromatic Heterocycles	0
Num Lipinski HB Donor	0
Num Atoms	23
Num Lipinski HB Acceptor	3
Num Rings	3

Potential Toxicity:

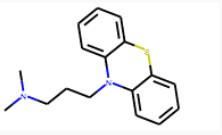
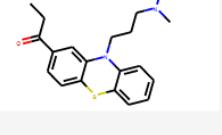
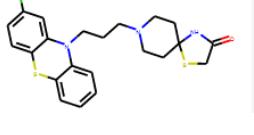
Index	Action	Injury	Confidence	Similar active compound in database
000	Modulator of Serotonin 2b (5-HT2b) receptor	Heart, Nervous system, Gastrointestinal	0.995	
001	Antagonist of the retinoic acid receptor (RAR) signaling pathway	Immune	0.981	
002	Disruptors of the mitochondrial membrane potential	Liver	0.928	

Figure S2. Example figure of result table from toxic substructure analysis.

Toxic substructure:		
Index	Alert Name	Structure
000	Hepatotoxicity	
001	Hepatotoxicity	
002	Alert from Top200 Drug	