## Support Information

## Table S1. Average Tanimoto similarity index in FHM, BS and RT data sets

	FHM	BS	RT
Total of compounds	995	900	903
Average Tanimoto similarity index	0.35	0.336	0.351

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		Methods	Model ID	Q	SE	SP	FP	FN
			1	0 (72	0.762	0.500	20	25
BS	Test (162)	4D-RF		0.075	0.762	0.509	28	25
		4D-k-NN	2	0.716	0.810	0.544	26	20
		FP4-RF	3	0.747	0.905	0.456	31	10
		FP4-SVM	4	0.778	0.905	0.544	26	10
FHM	Test (181)	4D-RF	5	0.751	0.621	0.872	12	33
		4D-k-NN	6	0.768	0.759	0.777	21	21
		FP4-RF	7	0.691	0.575	0.798	19	37
		FP4-SVM	8	0.696	0.690	0.702	28	27
RT	Test (162)	4D-RF	9	0.790	0.931	0.435	26	8
		4D-k-NN	10	0.747	0.836	0.522	19	22
		FP4-RF	11	0.809	0.957	0.435	26	5
		FP4-SVM	12	0.796	0.897	0.544	21	12
Global	Test (320)	4D-RF	13	0.728	0.683	0.791	28	59
		4D-k-NN	14	0.719	0.785	0.627	50	40
		FP4-RF	15	0.756	0.790	0.709	39	39
		FP4-SVM	16	0.772	0.780	0.761	32	41

**Table S2.** Using 4 descriptors and FP4 modeling results in binary classification of local models

		Methods	Model ID	Q	0-0	0-1	0-2	1-0	1-1	1-2	2-0	2-1	2-2
BS	Test (162)	4D-RF	a	0.586	45	1	11	20	3	13	21	1	47
		4D-k-NN	b	0.549	31	12	14	11	16	9	12	15	42
		FP4-RF	c	0.562	32	6	19	13	5	18	9	6	54
		FP4-SVM	d	0.599	37	7	13	14	10	12	11	8	50
FHM Test (		4D-RF	e	0.641	84	3	7	30	11	10	12	3	21
	$T_{out}(181)$	4D-k-NN	f	0.641	74	14	6	16	23	12	7	10	19
	Test (181)	FP4-RF	g	0.514	85	2	7	38	3	10	30	1	5
		FP4-SVM	h	0.558	78	12	4	27	10	14	15	8	13
RT 7	Test (162)	4D-RF	i	0.562	29	1	16	9	3	34	10	1	59
		4D-k-NN	j	0.568	28	7	11	12	21	13	9	18	43
		FP4-RF	k	0.617	35	3	8	15	9	22	11	3	56
		FP4-SVM	1	0.648	39	4	3	8	20	18	12	12	47

Table S3. The performance of ternary classification models in FHM, BS and RT

0-0 means non-toxicity predicted to non-toxicity, 0-1 means non-toxicity predicted to moderate-toxicity, 0-2 means nontoxicity predicted to high-toxicity (just like false positive in binary classification), 1-0 means moderate-toxicity predicted to non-toxicity, 1-1 means moderate-toxicity predicted to moderate-toxicity, 1-2 means moderate-toxicity predicted to high-toxicity, 2-0 means high-toxicity predicted to non-toxicity (just like FN in binary classification), 2-1 means hightoxicity predicted to moderate-toxicity, 2-2 means high-toxicity predicted to high-toxicity.



Table S4. The sensitive substructure alerts in FHM

## **Figure Captions**

Figure S1. Workflow of model building.

Figure S2. The tanimoto similarity index for the FHM, BS and RT's data set

**Figure S3.** The radar chart of five physicochemical descriptors: CrippenLogP, ATSm1, SwHBa and ETA\_dEpsilon\_D and molecular weight for the global model data set were presented. Each color line represents a compound.

Figure S4. Representation of receiver operating characteristics (ROC) curve for the validation set in model 2, 5, 10 and 14.



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