

Table S9: The accuracy and ROC-AUC value of five major CYP Isoforms from PubChem Data Set II. ImageMol_NonPretrained and ImageMol indicate that ImageMol is not pre-trained and ImageMol is pre-trained on 10M molecular images. "MACCS-" represents the method based on MACCS molecular fingerprint, "FP4-" represents the method based on FP4 molecular fingerprint. "CC-" represents combined algorithm (ensemble learning). For details on other methods, see [1]. ImageMol_NonPretrained and ImageMol are fine-tuned on PubChem Data Set I and evaluated on PubChem Data Set II. The best and second best result for each dataset is bolded and underlined, respectively. The methods in blue background represent image-based methods. The "3 times" represents the mean \pm standard deviation of 3 runs with different random seeds.

	ValidationSet									
	accuracy					ROC-AUC				
methods	1A2	2C9	2C19	2D6	3A4	1A2	2C9	2C19	2D6	3A4
MACCS-SVM	68.0	<u>86.6</u>	80.3	<u>87.8</u>	74.9	0.814	0.854	0.841	0.88	0.783
MACCS-C4.5 DT	68.3	78.1	77.5	82.6	70.6	–	–	–	–	–
MACCS-k-NN	69.0	81.5	80.7	84.5	70.4	–	–	–	–	–
MACCS-NB	61.0	69.4	70.5	80.6	69.3	–	–	–	–	–
MACCS-CC-I	71.3	86.5	80.6	87.5	75.1	0.809	0.857	0.838	0.88	0.78
MACCS-CC-II	72.0	86.7	80.8	87.5	74.9	0.806	<u>0.861</u>	0.842	0.88	0.777
MACCS-CC-III	72.3	86.3	80.5	87.5	74.6	0.814	0.85	0.839	0.872	0.775
MACCS-CC-IV	<u>73.1</u>	86.4	80.4	87.6	75.0	0.815	0.853	0.829	0.886	0.785
MACCS-CC-V	69.9	83.6	80.4	86.2	74.2	0.776	0.837	0.829	0.86	0.754
MACCS-Mean	68.1	84.0	76.6	87.2	73.6	0.801	0.845	0.756	<u>0.881</u>	0.778
MACCS-Maximum	67.0	81.5	79.7	86.0	72.8	0.789	0.813	0.67	0.874	0.765
MACCS-Multiply	67.1	82.7	79.5	86.8	73.7	0.795	0.829	0.704	0.879	0.772
FP4-SVM	67.6	84.2	80.5	86.4	75.6	0.803	0.832	0.82	0.848	0.785
FP4-C4.5 DT	65.8	78.8	77.2	78.5	72.0	–	–	–	–	–
FP4-k-NN	69.7	78.4	77.9	81.1	66.3	–	–	–	–	–
FP4-NB	59.7	75.4	73.5	80.2	70.9	–	–	–	–	–
FP4-CC-I	70.7	84.5	80.2	85.6	<u>75.8</u>	0.805	0.842	0.798	0.86	0.79
FP4-CC-II	71.2	84.6	81.0	86.1	<u>75.8</u>	0.798	0.85	0.815	0.86	0.783
FP4-CC-III	70.8	84.6	80.7	85.8	76.0	0.794	0.827	0.812	0.839	0.787
FP4-CC-IV	71.7	84.1	80.7	85.6	75.4	0.805	0.837	0.793	0.842	0.792
FP4-CC-V	70.2	83.2	79.0	84.5	75.0	0.764	0.844	0.819	0.841	0.777
FP4-Mean	67.2	81.9	77.5	85.6	74.8	0.782	0.848	0.774	0.856	<u>0.79</u>
FP4-Maximum	66.5	80.0	78.1	83.9	74.0	0.772	0.836	0.674	0.843	0.784
FP4-Multiply	67.3	81.1	77.9	85.2	73.9	0.778	0.842	0.729	0.851	0.788
Chemception	65.1	66.7	67.1	68.1	69.3	0.699	0.733	0.738	0.753	0.763
ADMET-CNN	62.5	73.3	76.4	81.0	70.5	0.556	0.697	0.616	0.559	0.654
QSAR-CNN	55.6	64.1	60.1	82.1	60.0	0.712	0.736	0.752	0.761	0.734
ImageMol_NonPretrained	73.3	84.3	<u>82.2</u>	85.7	74.2	<u>0.837</u>	0.841	<u>0.848</u>	0.83	0.763
ImageMol	71.9	85.0	82.9	88.3	74.6	0.852	0.870	0.871	0.893	0.799
Chemception (3 times)	64.8 \pm 0.4	67.1 \pm 0.4	65.9 \pm 1.2	67.5 \pm 0.7	69.2 \pm 0.2	0.690 \pm 0.010	0.727 \pm 0.005	0.728 \pm 0.007	0.744 \pm 0.09	0.761 \pm 0.002
ADMET-CNN (3 times)	62.2 \pm 0.4	76.4 \pm 0.0	73.3 \pm 0.0	81.1 \pm 0.1	70.5 \pm 0.0	0.546 \pm 0.008	0.668 \pm 0.020	0.603 \pm 0.009	0.565 \pm 0.005	0.652 \pm 0.003
QSAR-CNN (3 times)	55.9 \pm 2.0	64.1 \pm 0.1	59.9 \pm 0.4	81.6 \pm 0.8	58.8 \pm 1.7	0.710 \pm 0.003	0.734 \pm 0.009	0.735 \pm 0.007	0.759 \pm 0.008	0.734 \pm 0.003
ImageMol (3 times)	74.0\pm0.8	84.7\pm0.4	83.1\pm0.2	88.3\pm0.03	75.6\pm0.3	0.845\pm0.001	0.874\pm0.004	0.868\pm0.003	0.890\pm0.003	0.796\pm0.001

[1] Cheng F, Yu Y, Shen J, et al. Classification of cytochrome P450 inhibitors and noninhibitors using combined classifiers[J]. Journal of chemical information and modeling, 2011, 51(5): 996–1011.