

## Supplementary Tables and Figures

**Table S1.** The pharmacophore hypothesis for PDE4D ligand.

Hypothesis	Total Cost	Cost difference	Error Cost	RMS	R value	Feature
1	305.64	190.78	287.965	1.601	0.851	AHHHZ
2	308.29	188.13	292.176	1.643	0.842	AHHHH
3	311.09	185.33	295.102	1.671	0.836	AHHHH
4	311.30	185.13	295.178	1.672	0.836	AHHHH
5	312.62	183.81	296.513	1.685	0.833	AHHHH
6	313.01	183.41	294.947	1.669	0.837	AHHH
7	314.82	181.61	297.399	1.693	0.832	AHHHZ
8	316.28	180.15	299.873	1.716	0.826	AHHHH
9	316.54	179.89	298.248	1.701	0.830	AHHH
10	317.67	178.76	301.657	1.733	0.823	AHHHH

Null cost: 496.43. Fixed cost: 224.52. Configuration cost: 14.851. Hydrogen bond acceptors (A), hydrophobic (H), and negative ionizable features (Z) were identified and analyzed for modeling.

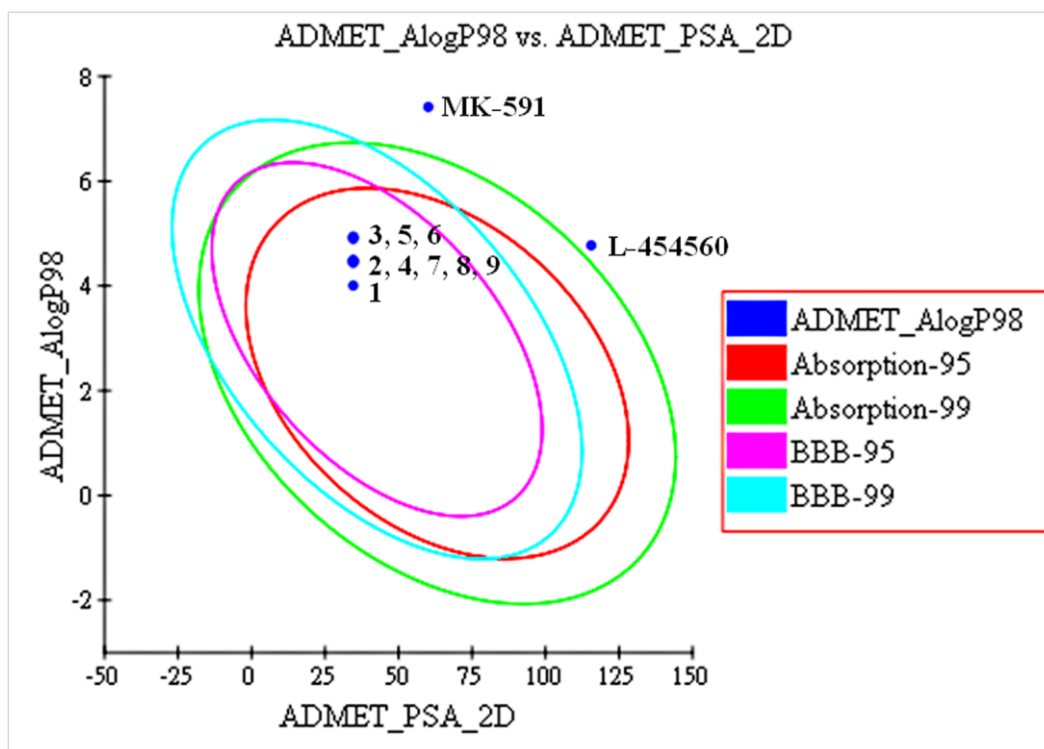
**Table S2.** The pharmacophore hypothesis for ALOX5AP ligand.

<b>Hypothesis</b>	<b>Total Cost</b>	<b>Cost difference</b>	<b>Error Cost</b>	<b>RMS</b>	<b>R value</b>	<b>Feature</b>
1	153.97	41.92	137.94	1.669	0.768	AHHHH
2	158.10	37.80	142.489	1.761	0.737	AHHHZ
3	162.57	33.33	146.511	1.838	0.709	AHHHZ
4	164.18	31.71	148.412	1.873	0.695	AHHHH
5	164.20	31.70	148.211	1.869	0.697	AHHHH
6	164.87	31.03	149.201	1.887	0.689	AHHHH
7	165.76	30.13	147.518	1.856	0.702	AHHHZ
8	166.20	29.70	150.449	1.910	0.680	AHHHH
9	166.72	29.17	149.387	1.891	0.688	AHHHH
10	166.82	29.08	151.152	1.923	0.675	AHHH

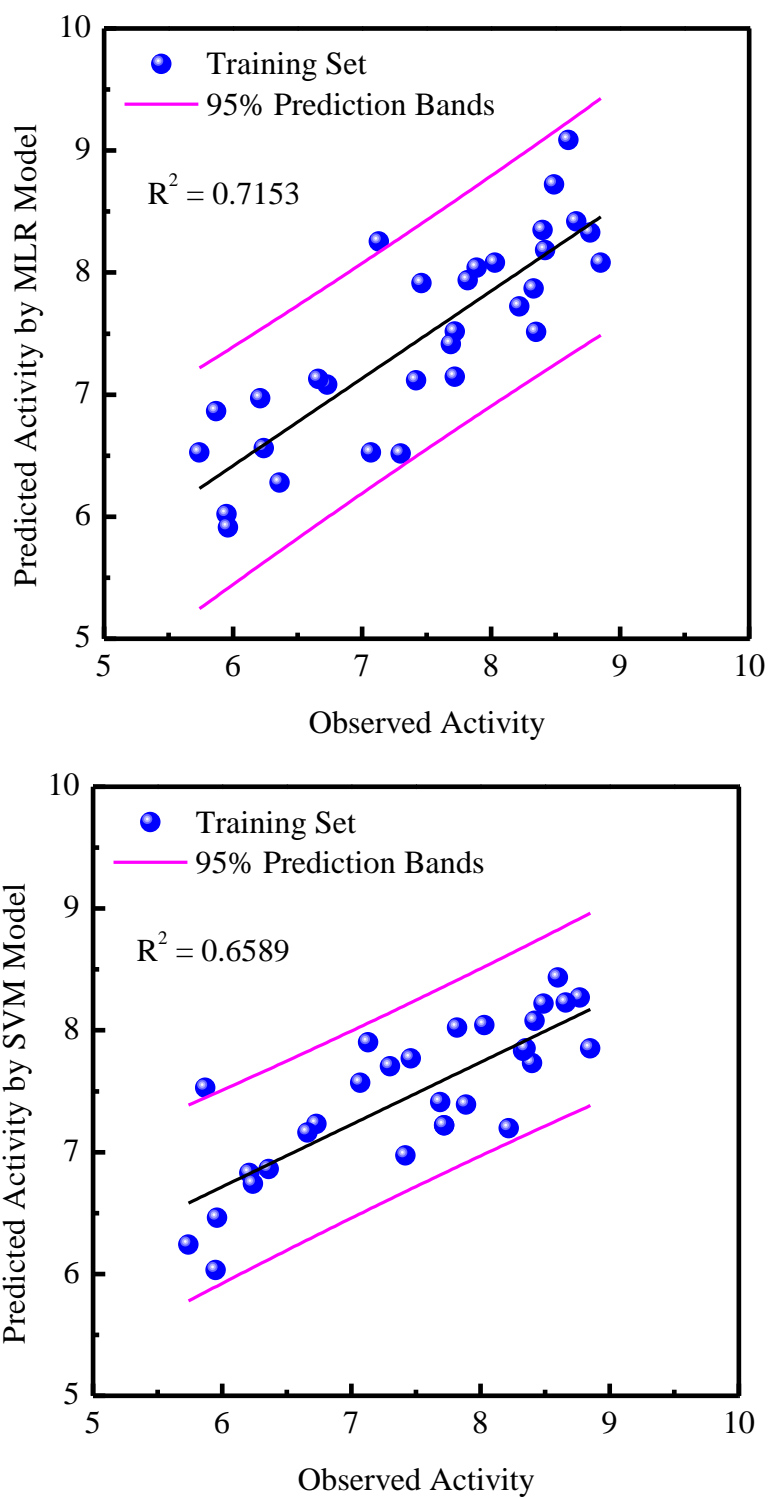
Null cost: 195.90. Fixed cost: 113.12. Configuration cost: 14.452. Hydrogen bond acceptors (A), hydrophobic (H), and negative ionizable features (Z) were identified and analyzed for modeling.

**Table S3.** The interaction energy (kcal/mol) with divalent metal cations for both TCM candidates and L-454560.

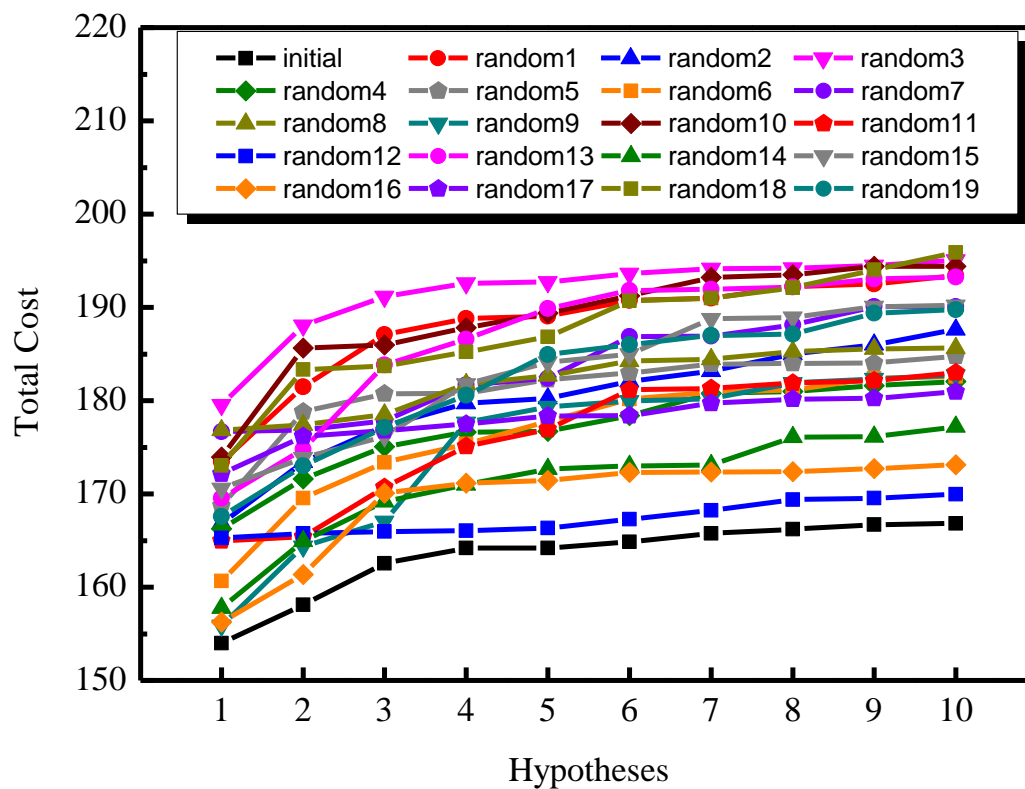
Ligand	$Zn^{2+}$			$Mg^{2+}$		
	Interaction Energy	Electrostatic Interaction Energy	VDW Interaction Energy	Interaction Energy	Electrostatic Interaction Energy	VDW Interaction Energy
Myristic acid	-290.095	-297.044	6.949	-254.020	-259.442	5.422
Pentadecanoic acid	-257.359	-262.071	4.713	-279.822	-284.465	4.643
L-454560	-80.551	-86.800	6.249	-16.621	-16.473	-0.147



**Figure S1.** Human intestinal absorption model for (1) Myristic acid, (2) Pentadecanoic acid, (3) 2-pentadecenoic acid, (4) 2-hexadecenoic acid, (5) 9,12-Octadecadienoic acid, (6) Hexadecanoic acid, (7) 11-hexadecenoic acid, (8) Palmitoleic acid, (9) Punicic acid, and controls, L-454560 and MK-591.



**Figure S2.** Comparative plots of MLR (top) and SVM (bottom) models for ALOX5AP inhibitors. Correlation trend (black line) and 95% confidence regions (enclosed by magenta lines) were shown. Training set (blue dots) was presented.



**Figure S3.** CatScramble validation plot for ALOX5AP that compares Total costs between initial and randomized HypoGen hypothesis.