Supporting Information

Insight into Shared Properties and Differential Dynamics and Specificity of Secretory Phospholipase A_2 Family Members

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This file contains: 13 pages, 5 figures and 2 tables.

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Supplementary Figures

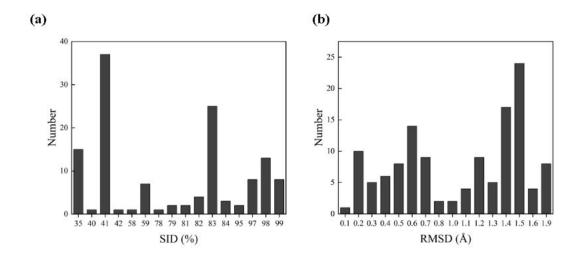


Figure S1. Distributions of percent sequence identity (SID) (a) and RMSDs (b) with respect to the reference structure 1vkq from group IB for the 130 sPLA_2 family members.

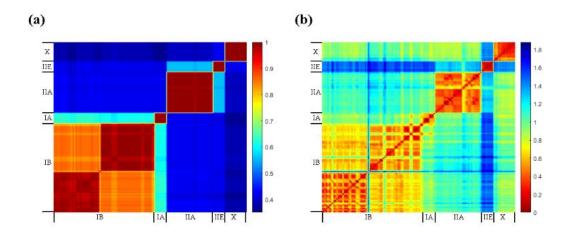


Figure S2. Heat maps of sequence identity (a) and RMSDs (b) between members for 130 sPLA₂ family members from groups IA, IB, IIA, IIE and X.

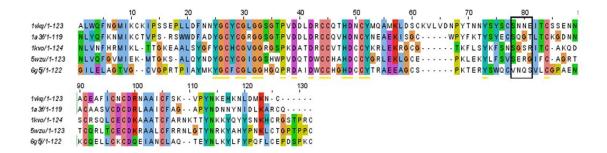


Figure S3. Diagram of sequence alignment of the five representative structures for groups IA, IB, IIA, IIE and X of $sPLA_2$ enzyme family. The sequences of region 3 are marked in the black box.

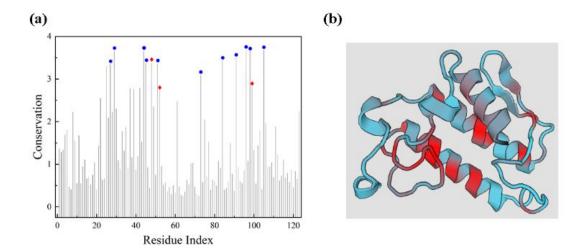


Figure S4. (a) Residue conservation of sPLA₂ family from a multiple sequence alignment of 789 sequences, with the sequence numbering corresponding to the reference structure 1vkq. The highly conserved residues Cys27, Cys29, Cys44, Cys45, Cys51, Cys84, Cys91, Cys96, Cys98, Cys105 and Tyr73 are marked in blue, and the other highly conserved residues His48, Tyr52 and Asp99 (catalytic residues) are marked in red. (b) Color-coded residue conservation of sPLA₂ enzyme mapped on the reference structure 1vkq (download from MISTIC web). A high level in conservation is shown in red and a low level is shown in blue.

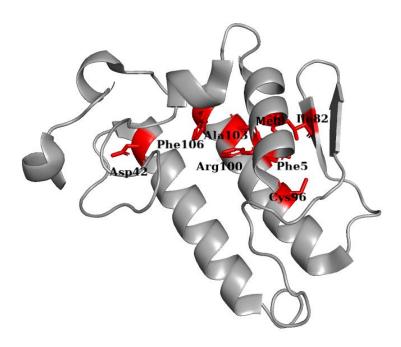


Figure S5. Residues (in red) with minimal sensitivity which are also characterized by the lowest variances among sPLA₂ enzyme members, are tightly packed in the structure.

Supplementary Tables

Table S1. The 130 sPLA_2 enzyme members used in this study.

	PDB	Uniprot ID	Protein Name	Sequence	RMSD (Å)
				identity with	from the
				the reference	reference
				one	one
Grou	p IB				
1	1vkq-A	PA21B BOVIN	phospholipase A ₂	1.000	0.000
2	1bpq-A	PA21B BOVIN	phospholipase A ₂	0.984	0.227
3	1bp2-A	PA21B BOVIN	phospholipase A ₂	0.976	0.441
4	1c74-A	PA21B BOVIN	phospholipase A ₂	0.992	0.197
5	1ceh-A	PA21B BOVIN	phospholipase A ₂	0.967	0.328
6	1fdk-A	PA21B BOVIN	phospholipase A ₂	0.976	0.259
7	1g4i-A	PA21B BOVIN	phospholipase A ₂	0.976	0.454
8	1gh4-A	PA21B BOVIN	phospholipase A ₂	0.984	0.830
9	1irb-A	PA21B BOVIN	phospholipase A ₂	0.967	0.246
10	1kvw-A	PA21B BOVIN	phospholipase A ₂	0.967	0.441
11	1kvx-A	PA21B BOVIN	phospholipase A ₂	0.967	0.466
12	1kvy-A	PA21B BOVIN	phospholipase A ₂	0.967	0.381
13	1mks-A	PA21B BOVIN	phospholipase A ₂	0.951	0.300
14	1mkt-A	PA21B BOVIN	phospholipase A ₂	0.976	0.262
15	1mku-A	PA21B BOVIN	phospholipase A ₂	0.951	0.562
16	1mkv-A	PA21B BOVIN	phospholipase A ₂	0.976	0.245
17	1o2e-A	PA21B BOVIN	phospholipase A ₂	1.000	0.241
18	1o3w-A	PA21B BOVIN	phospholipase A ₂	1.000	0.165
19	1une-A	PA21B BOVIN	phospholipase A ₂	0.976	0.441
20	1vl9-A	PA21B BOVIN	phospholipase A ₂	0.984	0.560
21	2b96-A	PA21B BOVIN	phospholipase A ₂	0.984	0.308
22	2bax-A	PA21B BOVIN	phospholipase A ₂	0.992	0.141
23	2bch-A	PA21B BOVIN	phospholipase A ₂	0.992	0.170
24	2bd1-A	PA21B BOVIN	phospholipase A ₂	0.992	0.607
25	2bd1-B	PA21B BOVIN	phospholipase A ₂	0.992	0.597
26	2bp2-A	PA21B BOVIN	phospholipase A ₂	0.976	0.531
27	2bpp-A	PA21B BOVIN	phospholipase A ₂	0.976	0.236
28	2zp3-A	PA21B BOVIN	phospholipase A ₂	0.967	0.204
29	2zp4-A	PA21B BOVIN	phospholipase A ₂	0.967	0.199
30	2zp5-A	PA21B BOVIN	phospholipase A ₂	0.967	0.360
31	3bp2-A	PA21B BOVIN	phospholipase A ₂	0.975	0.502
32	3elo-A	PA21B HUMAN	phospholipase A ₂	0.780	1.485

33	1fx9-A	PA21B PIG	Phospholipase A ₂ ,	0.829	0.665
			major isoenzyme		
34	1fx9-B	PA21B PIG	Phospholipase A ₂ ,	0.829	0.661
			major isoenzyme		
35	1fxf-A	PA21B PIG	Phospholipase A ₂ ,	0.829	0.633
			major isoenzyme		
36	1fxf-B	PA21B PIG	Phospholipase A ₂ ,	0.829	0.624
			major isoenzyme		
37	118s-A	PA21B PIG	Phospholipase A ₂ ,	0.829	0.642
			major isoenzyme		
38	118s-B	PA21B PIG	Phospholipase A ₂ ,	0.829	0.625
			major isoenzyme		
39	1p2p-A	PA21B PIG	Phospholipase A ₂ ,	0.829	0.604
			major isoenzyme		
40	1y6o-A	PA21B PIG	Phospholipase A ₂ ,	0.821	0.450
			major isoenzyme		
41	1y6o-B	PA21B PIG	Phospholipase A ₂ ,	0.821	0.435
			major isoenzyme		
42	1y6p-A	PA21B PIG	Phospholipase A ₂ ,	0.821	0.754
			major isoenzyme		
43	1y6p-B	PA21B PIG	Phospholipase A ₂ ,	0.821	1.037
			major isoenzyme		
44	2azy-A	PA21B PIG	Phospholipase A ₂ ,	0.829	1.209
			major isoenzyme		
45	2azz-A	PA21B PIG	Phospholipase A ₂ ,	0.829	1.242
			major isoenzyme		
46	2b00-A	PA21B PIG	Phospholipase A ₂ ,	0.829	1.212
			major isoenzyme		
47	2b01-A	PA21B PIG	Phospholipase A ₂ ,	0.829	1.041
			major isoenzyme		
48	2b03-A	PA21B PIG	Phospholipase A ₂ ,	0.829	1.307
			major isoenzyme		
49	2b04-A	PA21B PIG	Phospholipase A ₂ ,	0.838	1.385
			major isoenzyme		
50	2phi-A	PA21B PIG	Phospholipase A ₂ ,	0.837	0.569
			major isoenzyme		
51	2phi-B	PA21B PIG	Phospholipase A ₂ ,	0.837	0.505
			major isoenzyme		
52	3fvi-A	PA21B PIG	Phospholipase A ₂ ,	0.829	0.563
			major isoenzyme		
53	3fvi-B	PA21B PIG	Phospholipase A ₂ ,	0.829	0.484
			major isoenzyme		

54	3fvi-C	PA21B PIG	Phospholipase A ₂ ,	0.829	0.499
	20:5	D. O.I.D. DV.C.	major isoenzyme	0.020	0.500
55	3fvi-D	PA21B PIG	Phospholipase A ₂ ,	0.829	0.560
			major isoenzyme		
56	3fvj-A	PA21B PIG	Phospholipase A ₂ ,	0.829	0.690
			major isoenzyme		
57	3hsw-A	PA21B PIG	Phospholipase A ₂ ,	0.829	0.691
			major isoenzyme		
58	3130-A	PA21B PIG	Phospholipase A ₂ ,	0.829	0.632
			major isoenzyme		
59	3o4m-A	PA21B PIG	Phospholipase A ₂ ,	0.829	0.661
			major isoenzyme		
60	3p2p-A	PA21B PIG	Phospholipase A ₂ ,	0.805	1.139
			major isoenzyme		
61	3p2p-B	PA21B PIG	Phospholipase A ₂ ,	0.805	1.097
			major isoenzyme		
62	3qlm-A	PA21B PIG	Phospholipase A ₂ ,	0.829	0.676
			major isoenzyme		
63	4dbk-A	PA21B PIG	Phospholipase A ₂ ,	0.829	0.662
			major isoenzyme		
64	4g5i-A	PA21B PIG	Phospholipase A ₂ ,	0.829	0.620
			major isoenzyme		
65	4o1y-A	PA21B PIG	Phospholipase A ₂ ,	0.829	0.696
			major isoenzyme		
66	4p2p-A	PA21B PIG	Phospholipase A ₂ ,	0.829	0.650
			major isoenzyme		
67	5p2p-A	PA21B PIG	Phospholipase A ₂ ,	0.788	1.123
			major isoenzyme		
68	5p2p-B	PA21B PIG	Phospholipase A ₂ ,	0.788	1.093
			major isoenzyme		
Grou	p IA				
69	1a3f-A	PA2A2 NAJNA	Acidic phospholipase	0.590	1.215
			$A_2 2$		
70	1a3d-A	PA2A2 NAJNA	Acidic phospholipase	0.590	1.268
			$A_2 2$		
71	1a3f-B	PA2A2 NAJNA	Acidic phospholipase	0.590	1.222
			$A_2 2$		
72	1a3f-C	PA2A2 NAJNA	Acidic phospholipase	0.590	1.219
			A_2 2		
73	1psh-A	PA2A2 NAJNA	Acidic phospholipase	0.590	1.210
	•		A_2 2		
74	1psh-B	PA2A2 NAJNA	Acidic phospholipase	0.590	1.197

75	1psh-C	PA2A2 NAJNA	Acidic phospholipase A ₂ 2	0.590	1.215
76	2wq5-A	PA2A2 NAJNA	Acidic phospholipase A ₂ 2	0.581	1.303
Group	o IIA	1	'		
77	1kvo-A	PA2GA HUMAN	Phospholipase A ₂ , membrane associated	0.409	1.474
78	1ayp-A	PA2GA HUMAN	Phospholipase A ₂ , membrane associated	0.409	1.468
79	1ayp-B	PA2GA HUMAN	Phospholipase A ₂ , membrane associated	0.409	1.493
80	1ayp-C	PA2GA HUMAN	Phospholipase A ₂ , membrane associated	0.409	1.473
81	1ayp-D	PA2GA HUMAN	Phospholipase A ₂ , membrane associated	0.409	1.483
82	1ayp-E	PA2GA HUMAN	Phospholipase A ₂ , membrane associated	0.409	1.486
83	1ayp-F	PA2GA HUMAN	Phospholipase A ₂ , membrane associated	0.409	1.485
84	1db4-A	PA2GA HUMAN	Phospholipase A ₂ , membrane associated	0.409	1.427
85	1db5-A	PA2GA HUMAN	Phospholipase A ₂ , membrane associated	0.409	1.430
86	1dcy-A	PA2GA HUMAN	Phospholipase A ₂ , membrane associated	0.409	1.430
87	lj1a-A	PA2GA HUMAN	Phospholipase A ₂ , membrane associated	0.409	1.468
88	1j1a-B	PA2GA HUMAN	Phospholipase A ₂ , membrane associated	0.409	1.465
89	1kqu-A	PA2GA HUMAN	Phospholipase A ₂ , membrane associated	0.409	1.429
90	1kvo-B	PA2GA HUMAN	Phospholipase A ₂ , membrane associated	0.409	1.476
91	1kvo-C	PA2GA HUMAN	Phospholipase A ₂ , membrane associated	0.409	1.478
92	1kvo-D	PA2GA HUMAN	Phospholipase A ₂ , membrane associated	0.409	1.497
93	1kvo-E	PA2GA HUMAN	Phospholipase A ₂ , membrane associated	0.409	1.483
94	1kvo-F	PA2GA HUMAN	Phospholipase A ₂ , membrane associated	0.409	1.461
95	1n28-A	PA2GA HUMAN	Phospholipase A ₂ , membrane associated	0.409	1.496

96	1n28-B	PA2GA	Phospholipase A ₂ ,	0.409	1.557
		HUMAN	membrane associated		
97	1n29-A	PA2GA	Phospholipase A ₂ ,	0.417	1.525
		HUMAN	membrane associated		
98	1pod-A	PA2GA	Phospholipase A ₂ ,	0.409	1.583
		HUMAN	membrane associated		
99	1poe-A	PA2GA	Phospholipase A ₂ ,	0.409	1.595
		HUMAN	membrane associated		
100	1poe-B	PA2GA	Phospholipase A ₂ ,	0.409	1.503
		HUMAN	membrane associated		
101	3u8b-A	PA2GA	Phospholipase A ₂ ,	0.409	1.507
		HUMAN	membrane associated		
102	3u8d-A	PA2GA	Phospholipase A ₂ ,	0.409	1.422
		HUMAN	membrane associated		
103	3u8d-B	PA2GA	Phospholipase A ₂ ,	0.409	1.424
		HUMAN	membrane associated		
104	3u8h-A	PA2GA	Phospholipase A ₂ ,	0.409	1.470
		HUMAN	membrane associated		
105	3u8h-B	PA2GA	Phospholipase A ₂ ,	0.409	1.458
		HUMAN	membrane associated		
106	5g3n-A	PA2GA	Phospholipase A ₂ ,	0.409	1.461
		HUMAN	membrane associated		
107	5g3n-B	PA2GA	Phospholipase A ₂ ,	0.409	1.649
		HUMAN	membrane associated		
Grou	p IIE				•
108	5wzu-A	PA2GE	Phospholipase A ₂ ,	0.412	1.885
		HUMAN	membrane associated		
109	5wzm-A	PA2GE	Phospholipase A ₂ ,	0.411	1.887
		HUMAN	membrane associated		
110	5wzo-A	PA2GE	Phospholipase A ₂ ,	0.412	1.938
		HUMAN	membrane associated		
111	5wzs-A	PA2GE	Phospholipase A ₂ ,	0.412	1.878
		HUMAN	membrane associated		
112	5wzt-A	PA2GE	Phospholipase A ₂ ,	0.412	1.852
		HUMAN	membrane associated		
113	5wzv-A	PA2GE	Phospholipase A ₂ ,	0.412	1.874
		HUMAN	membrane associated		
114	5wzw-A	PA2GE	Phospholipase A ₂ ,	0.412	1.883
		HUMAN	membrane associated		
115	5y5e-A	PA2GE	Phospholipase A ₂ ,	0.402	1.899
	_	HUMAN	membrane associated		
		HUMAN	memorane associated		

116	6g5j-A	PA2GX	Group 10 secretory	0.348	1.357
		HUMAN	phospholipase A ₂		
117	1le6-A	PA2GX	Group 10 secretory	0.348	1.441
		HUMAN	phospholipase A ₂		
118	1le6-B	PA2GX	Group 10 secretory	0.348	1.506
		HUMAN	phospholipase A ₂		
119	1le6-C	PA2GX	Group 10 secretory	0.348	1.480
		HUMAN	phospholipase A ₂		
120	1le7-A	PA2GX	Group 10 secretory	0.348	1.355
		HUMAN	phospholipase A ₂		
121	1le7-B	PA2GX	Group 10 secretory	0.348	1.385
		HUMAN	phospholipase A ₂		
122	4uy1-A	PA2GX	Group 10 secretory	0.348	1.332
		HUMAN	phospholipase A ₂		
123	4uy1-B	PA2GX	Group 10 secretory	0.348	1.352
		HUMAN	phospholipase A ₂		
124	5g3m-A	PA2GX	Group 10 secretory	0.348	1.346
		HUMAN	phospholipase A ₂		
125	5g3m-B	PA2GX	Group 10 secretory	0.348	1.367
		HUMAN	phospholipase A ₂		
126	5ow8-A	PA2GX	Group 10 secretory	0.348	1.358
		HUMAN	phospholipase A ₂		
127	5ow8-B	PA2GX	Group 10 secretory	0.348	1.384
		HUMAN	phospholipase A ₂		
128	5owc-A	PA2GX	Group 10 secretory	0.348	1.350
		HUMAN	phospholipase A ₂		
129	5owc-B	PA2GX	Group 10 secretory	0.348	1.361
		HUMAN	phospholipase A ₂		
130	6g5j-B	PA2GX	Group 10 secretory	0.348	1.379
		HUMAN	phospholipase A ₂		

Table S2. The first ten pairs of residues ranked based on the mutual information between residues.

Residue	Residue	MI value
Cys11	Cys77	19.115
Tyr52	Met53	18.099
Ala102	Phe106	17.196
Phe5	Met8	14.956
Phe5	Ala102	14.593
Phe5	Ile9	14.041
Gly26	Gly33	13.591
His115	Leu118	13.363
Tyr111	Asn112	12.714
Ile13	Pro14	12.641