## **Supplementary information**

## Investigating the Impact of Asp181 Point Mutations on Interactions between PTP1B and Phosphotyrosine Substrate

Mengyuan Liu<sup>1</sup>, Lushan Wang<sup>2</sup>, Xun Sun<sup>1, \*</sup>, Xian Zhao<sup>1, \*</sup>

<sup>1</sup> State Key Laboratory of Crystal Materials, Shandong University, Jinan 250100, China,

<sup>2</sup> State Key Laboratory of Microbial Technology, Shandong University, Jinan 250100, China.

Correspondence and requests for materials should be addressed to X. S. (sunxun@icm.sdu.edu.cn)

and X. Z. (zhaoxian@icm.sdu.edu.cn)



**Figure S1** | **Comparison of RMSD values in different regions.** (A) P loop. (B) WPD loop. (C) Secondary aryl-phosphate-binding site. (D) Other residues in the active site of PTP1B.

Complex	Donor	Acceptor	Distribution <sup>a</sup>
WT	S216-N-H	PTR-O3P	xxx oxxx*xxoxxxxxxxxxxx***xooxxxooxxxox
	A217-N-H	PTR-O1P	*000000*.
	A217-N-H	PTR-O3P	*xxx**xxxxxx**xx*xxx*xxxxxxxxxxxx
	G218-N-H	PTR-O3P	000-000
	I219-N-H	PTR-O2P	@0x *@@@@@@@@@@@@@@@@@@@@@@@@@@@@@@@@@@@
	I219-N-H	PTR-O3P	*000000.
	G220-N-H	PTR-O2P	@0x*@0**@0*00@0@0*@00**@0***@*x@0@0**@0@0
	G220-N-H	PTR-O3P	*@@@@@@.
	R221-N-H	PTR-O1P	00x *0000000000000000000000000000000000
	R221-N-H	PTR-O2P	*000000.
	R221-NE-HE	PTR-O1P	@@x -oo-xoo*@@**@@@@@@@@@@@@@@@@@@@#@*@@@@@
	R221-NE-HE	PTR-O2P	00000000.
	R221-NH2-HH21	PTR-O1P	*000000
	R221-NH2-HH21	PTR-O3P	@@* *@@@@@@@@@@@@@@@@@@@@@@@@@@@@@@@@@
D181A	S216-N-H	PTR-O3P	x-0-xx00000000000 00
	A217-N-H	PTR-O2P	
	A217-N-H	PTR-O3P	@@@@*@@**@@@*****@@@@@@@@@@@@
	G218-N-H	PTR-O2P	*************X*X*@**@X**X*X OXX***XX**
	I219-N-H	PTR-O1P	-@@@@@@@
	I219-N-H	PTR-O2P	00000000000000000000000000000000000000
	G220-N-H	PTR-O1P	·
	G220-N-H	PTR-O2P	@@@@@@@@@@@@@@@@@@@@@@@@@@@@@@@@@@@@@@
	R221-N-H	PTR-O1P	00000*00000000000000000000000000000000
	R221-N-H	PTR-O3P	-@*@@@@@
	R221-NE-HE	PTR-O1P	@@@@@@@@@@@@@@@@@@@@@@@@@@@@@@@@@@@@@@
	R221-NE-HE	PTR-O2P	@@@@-
	R221-NH2-HH21	PTR-O1P	0
	R221-NH2-HH21	PTR-O3P	@@@@@@@@@@@@@@@@@@@@@@x X@@@@@@@@@
D181E	S216-N-H	PTR-O3P	XXX*X***XX**X****@@*****X*@***@***@***@
	A217-N-H	PTR-O3P	@@*@@@*@*@@@#**@ <sub>X</sub> @* <sub>X</sub> *****@*@*@* <sub>X</sub> ***** <sub>X</sub> *** <sub>X</sub> *** <sub>X</sub> ******
	G218-N-H	PTR-O2P	
	I219-N-H	PTR-O2P	000000000000000000000000000000000000000
	G220-N-H	PTR-O2P	02
	R221-N-H	PTR-O1P	02
	R221-NE-HE	PTR-O1P	02
	R221-NH2-HH21	PTR-O1P	
	R221-NH2-HH21	PTR-O3P	000000000000000000000000000000000000000

Table S1 The analysis of hydrogen bonds between PTP1B and substrate during the last 30 ns MD simulations.

<sup>a</sup> Hydrogen bond information dumped for occupancies with dumping schematic of time series (every 60 ps) after each H-bond, key follows: "" 0-5%; "." 5-20%; "-" 20-40%; "o": 40-60%; "x": 60-80%; "\*": 80-95%; "@": 95-100%



Figure S2 | The average of electrostatic interaction energies between substrate and residues in the active site of PTP1B.



Figure S3 | The average of van der Waals interaction energies between substrate and residues in the active site of PTP1B.