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Figure S1. Pharmacophore models considered in this study. A) pharmacophore hypothesis generated from the glyphosate-EPSPS interactions (PDB ID: 3FJZ); B) pharmacophore model based on the overlay of multiple active conformations of glyphosate (PDB IDs: 3FJZ, 2QFU, 2GGA, 2AAY, 1G6S, 3NVS, 3SLH, 1RF6); C) pharmacophore hypothesis built from the intermediate state-like inhibitor (ISLI)-EPSPS interactions (PDB ID: 2PQC); D) pharmacophore model based on the overlay of multiple active conformations of ISLI (PDB IDs: 2PQC, 2PQD). Light blue sphere indicates H-bond donors, light red – H-bond acceptor, blue – positive ionic group, red – negative ionic group. All models were visualized in Schrodinger Maestro 12.6.



Figure S2. ROC curves built for the different pharmacophore models to estimate their screening performance and make prediction about their future performance in actual screening. A) and B) ROC curves for the glyphosate-interaction-based pharmacophore screened against DUDE and RADER validation datasets, respectively; C) and D) ROC curves for the glyphosate active conformations-based pharmacophore screened against DUDE and RADER validation datasets, respectively; E) and F) ROC curves for the ISLI-interaction-based pharmacophore screened against DUDE and RADER validation datasets, respectively; G) and H) ROC curves for the ISLI active conformations-based pharmacophore screened against DUDE and RADER validation datasets, respectively; G) and H) ROC curves for the ISLI active conformations-based pharmacophore screened against DUDE and RADER validation datasets, respectively; G) and H) ROC curves for the ISLI active conformations-based pharmacophore screened against DUDE and RADER validation datasets, respectively; G) and H) ROC curves for the ISLI active conformations-based pharmacophore screened against DUDE and RADER validation datasets, respectively; G) and H) ROC curves for the ISLI active conformations-based pharmacophore screened against DUDE and RADER validation datasets, respectively.

Number	ChEMBL id	Name	Inhibitory potency (K _i)
1	CHEMBL95406	(N-Phosphonomethyl-hydrazino)-acetic acid	0.61 µM
2	CHEMBL96010	N-Nitroso-N-(phosphonomethyl)glycine	2.20 µM
3	CHEMBL97331	2-{3-carboxy-5- [carboxy(phosphono)methoxy]phenoxy}propanedioic acid	1.30 µM
4	CHEMBL97756	2-[methyl(phosphonomethyl)amino]acetic acid	78 µM
5	CHEMBL98618	{[(2-methoxy-2-oxoethyl)amino]methyl}phosphonic acid	1.00 µM
6	CHEMBL98868 (CHEMBL95764)	2-[(phosphonomethyl)amino]acetic acid (glyphosate)	0.16-52 μM
7	CHEMBL1627013 (CHEMBL1159978)	(3R,4S,5R)-5-[(1S)-1-carboxy-1- phosphonoethoxy]-4-hydroxy-3- (phosphonooxy)cyclohex-1-ene-1-carboxylic acid	0.11-5 nM
8	CHEMBL609383	(3R,4S,5R)-5- [(carboxymethyl)(phosphonomethyl)amino]-4-hydrox y-3- (phosphonooxy)cvclohex-1-ene-1-carboxylic acid	7.4-13.0 μM

9	CHEMBL1627015 (CHEMBL1206855)	(3R,4S,5R)-5-[(1S)-1-carboxy-2,2-difluoro-1- (phosphonooxy)ethoxy]-4-hydroxy-3- (phosphonooxy)cyclohex-1-ene-1-carboxylic acid	4 nM			
10	CHEMBL337302	(2E)-6,7-dihydroxy-2-[(2-hydroxy-4- methoxyphenyl)methylidene]-1-benzofuran-3-one	0.10 µM			
11	CHEMBL334500	(2E)-2-{[4-(diethylamino)phenyl]methylidene}-6,7- dihydroxy-1-benzofuran-3-one	0.65 µM			
12	CHEMBL330146	2-[hydroxy(phosphonomethyl)amino]acetic acid	0.65 µM			
12	CHEMBL329115	2-[3-carboxy-5-	2.50 µM			
15		(dicarboxymethoxy)phenoxy]propanedioic acid				
1/	CHEMBI 318618	(3R,4S,5R)-5-(carboxymethoxy)-4-hydroxy-3-	1.50 uM			
	CHEMDESTOOTO	(phosphonooxy)cyclohex-1-ene-1-carboxylic acid	1.50 μινι			
		3-[carboxylato(hydrogen				
15	CHEMBL313253	phosphonato)methoxy]-5-(hydrogen	0.16 µM			
		phosphonatooxy)benzoic acid				
		(3R,4S,5R)-5-(carboxylatomethoxy)-3-				
16	CHEMBI 287334	(hydrogen	1.50 uM			
10	ONEMBEZOVOOT	phosphonatooxy)-4-hydroxycyclohex-1-ene-1-carbo	1.00 pm			
		xylic acid				
17	CHEMBL1627011	3-[carboxy(phosphono)methoxy]-5-	0.16 µM			
	(CHEMBL1205445)	(phosphonooxy)benzoic acid				
18	CHEMBL1206828	3,5-bis(phosphonooxy)benzoic acid	0.78 µM			
19	CHEMBL319218	2-{[(1R,5R,6S)-3-carboxy-5-(carboxyformamido)-6-h	5.20 µM			
		ydroxycyclohex-2-en-1-yljoxy}propanedioic acid				
20	CHEMBL314765	3-carboxy-5-(hydrogen	0.78 uM			
		phosphonatooxy)phenyl hydrogen phosphate				
	CHEMBL1205418	(3R,4S,5R)-5-[(1S)-1-carboxy-2,2,2-trifluoro-1-	00 M			
21	(CHEMBL1627010)	(phosphonooxy)ethoxy]-4-hydroxy-3-	32 nM			
		(phosphonooxy)cyclohex-1-ene-1-carboxylic acid				
22	CHEMBL1205355	(3R,4R,5R)-5-(carboxymethoxy)-4-hydroxy-3-	1.50 µM			
	(CHEMBL88396)	(pnosphonooxy)cyclonex-1-ene-1-carboxylic acid	· ·			
23	CHEMBL1160691	4-[nydroxy(pnospnono)metnyi]-1H-pyrrole-2-	14.5 nM			
04		(3R,45)-4-nyaroxy-3-(pnospnonooxy)cyclonex-1-en	40			
24	CHEMBL1160690	e-I-	13 µM			
25	CHEMBL88396	(3K,4K,3K)-3-(CAIDOXYIAIOMEINOXY)-3-	1.5 µM			
		(nyurogen				
		phosphonalooxy -4-hydroxycyclonex- i-ene- i-carbo				
	Table S1 list of compounds from ChEMRI databases with domonstrated					
inhibitory potency against EDSDS any manufact to concrete the validation						
detecte						

datasets. *Remark*: all K_i values are for *E. coli* K12 EPSPS (id of the target – CHEMBL5033).



Figure S3. RMSD plots of all ligand atoms in the WT *Ei*EPSPS (red) and its poses in 50 ns, 100 ns, 115 ns and 150 ns analyzed over 150-ns MD simulation.



Figure S4. RMSD plots of all ligand atoms in the MUT *Ei*EPSPS (red) and its poses in 50 ns, 100 ns, 115 ns and 150 ns analyzed over 150-ns MD simulation.



Figure S5. The 2D diagram of the A) $EiEPSPS_{wt}$ and B) $EiEPSPS_{mut}$ complexed with AG351308, showing the average distances of the main intermolecular interactions with occupancy value $\ge 50\%$ obtained over the MD simulation.



Figure S6. RMSD plots of the backbone atoms in the WT and MUT ligand complexes (black) and only ligands (red) analyzed over 150 ns of MD simulation.



Figure S7. The 2D diagram of the A) $EiEPSPS_{wt}$ and B) $EiEPSPS_{mut}$ complexed with Glyphosate, showing the average distances of the main intermolecular interactions with occupancy value $\geq 50\%$ obtained over the MD simulation.



Figure S8. Per-residue energy decomposition analysis of main residues with energy decomposition value \leq or \geq 0.5 kcal/mol of of A) Glyphosate and B) AG332841 complexed with *Ei*EPSPS_{wt} (yellow) and *Ei*EPSPS_{mut} (black) structures.