

S1 Table. Comparison of the effector-binding sites (pockets) from the crystal structures of ligand-free DasR-EBD, ligand-free DasR and effector-bound DasR-EBD. The pocket volumes were calculated as described in the ‘Materials and Methods’ section.

Structure	ligand-free DasR-EBD	ligand-free DasR		DasR-EBD + GlcN-6-P		DasR-EBD + GlcNAc-6-P	
	A (-)	A (-)	B (-)	A (α)	B (β)	A (α)	B (β)
chain-ID (effector configuration)	A (-)	A (-)	B (-)	A (α)	B (β)	A (α)	B (β)
MS volume ^a (\AA^3)	1029.5	1126.8	1084.8	576.6	668.3	577.9	693.1
SA volume ^b (\AA^3)	218.0	252.0	271.3	74.7	94.1	94.9	101.8
pocket MS area ^c (\AA^2)	756.5	874.8	804.0	577.6	633.7	488.7	643.0
pocket SA area ^d (\AA^2)	426.8	456.1	422.1	196.5	238.0	217.8	251.1
# openings ^e	2	2	2	0	1	0	1
mouth MS area ^f (\AA^2)	146.9	157.3	160.6	-	12.1	-	28.1
mouth SA area ^g (\AA^2)	33.5	51.0	58.4	-	0.3	-	4.6
MS circumference ^h (\AA)	89.8	86.0	83.9	-	12.7	-	22.1
SA circumference ⁱ (\AA)	73.1	67.0	65.5	-	4.7	-	13.7

pocket volume based on the ^a molecular surface / ^b solvent-accessible surface
^c pocket molecular surface area / ^d pocket solvent-accessible surface area
^e number of mouths or openings to the external molecular surface
total area of mouth opening(s) based on the ^f molecular surface / ^g solvent-accessible surface
total circumference of mouth opening(s) based on the ^h molecular surface / ⁱ solvent-accessible surface