

S2 Table. DALI Statistics for structural alignment of AVR-Pia, AVR1-CO39, AVRPiz-t and ToxB.

RMSD (Å)	AVR-Pia	AVR1-CO39	AVRPiz-t	ToxB
Z-score				
AVR-Pia		2.3	2.8	2.2
AVR1-CO39	2.9		3.0	2.2
AVRPiz-t	3.3	3.1		2.2
ToxB	4.4	4.2	5.4	

The rmsd (upper right) and the Z-score (lower left) are from the pairwise superposition. The low RMSDs and the high Z-scores indicate that the ToxB(b) structure is the closest to the other three structures.