

Fig. S7 Overall structural comparison among representative type V effectors.

- (a) The overall structural similarity between $Cas\pi$ -1 and reported structures. The 10 structures with top Z-scores of similarities to $Cas\pi$ -1 are listed.
- (b) Structural similarity of protein domains within $Cas\pi-1$ to all reported structures. The reported structure with the highest Z-score of similarity to each $Cas\pi-1$ domain was shown.
- (c) The structural similarity between each $Cas\pi$ -1 domain and DpbCasX.
- (d) The structural similarity between each $Cas\pi$ -1 domain and LbCas12a.
- (e) The structures of representative type V effectors. All models were shown from the top view with the same size scale. For $Cas\pi$ and CasX, the domain organization within the primary sequence were shown. The protein domains within each structure are colored and labeled accordingly.