

Supplementary information, Figure S7

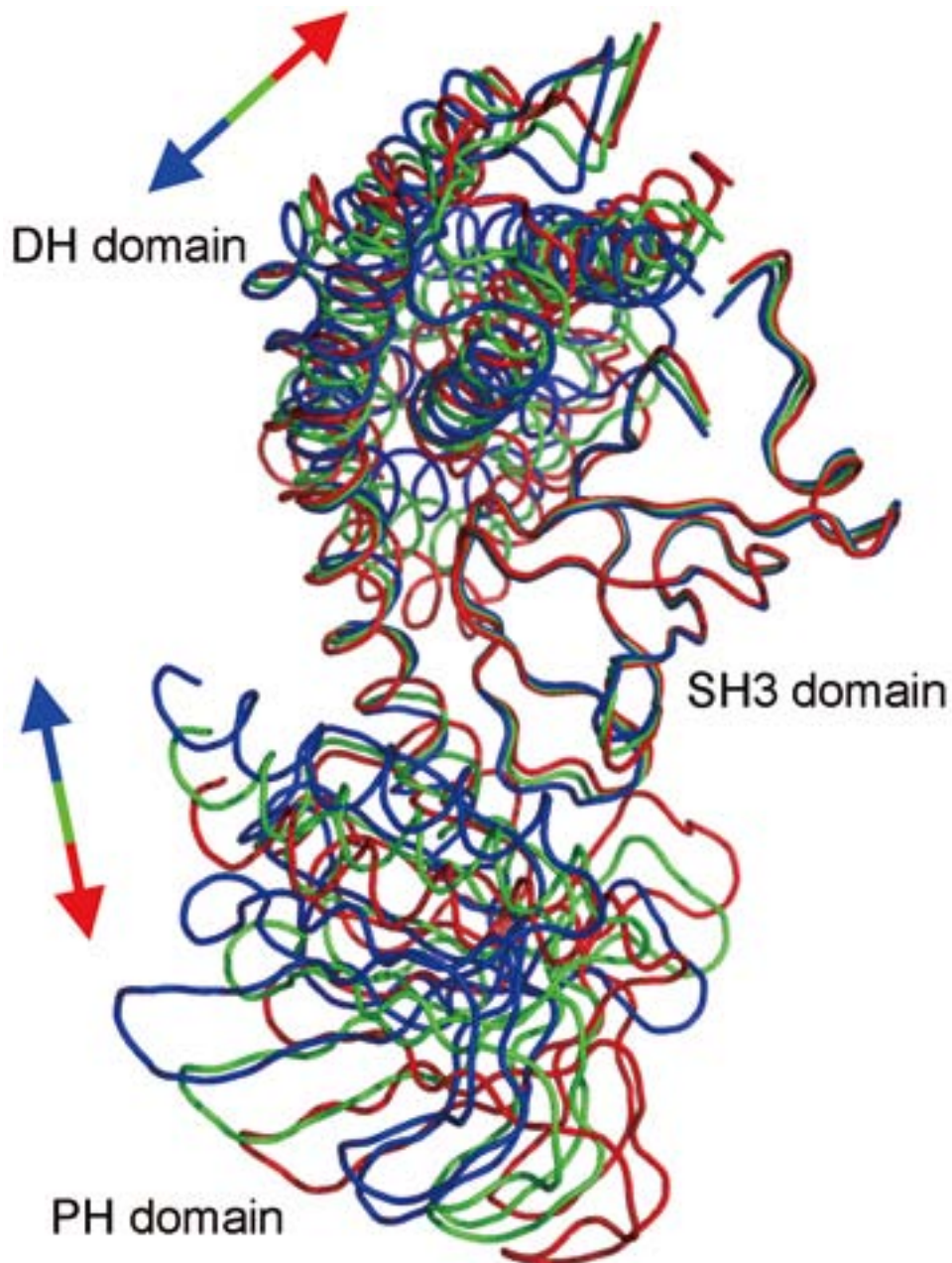


Figure S7 Normal mode analysis for the Asef protein. The crystal structure of autoinhibited Asef (PDB code: 2PZ1) was used as the initial state of Asef. For the 7th normal mode (the first six modes are translational and rotational modes), states 1, 8 and 24 (colored in green, red, and blue) of Asef from the normal mode analysis were superimposed by aligning their SH3 domains.