Supporting Information

Path Ensembles for Pin1-catalyzed Cis-trans Isomerization of a Substrate Calculated by Weighted Ensemble Simulations

Kei Moritsugu¹, Norifumi Yamamoto², Yasushige Yonezawa³, Shin-ichi Tate⁴, and Hiroshi Fujisaki^{5,6}*

¹Graduate School of Medical Life Science, Yokohama City University, 1-7-29 Suehirocho, Tsurumi, Yokohama 230-0045, Japan

²Department of Applied Chemistry, Faculty of Engineering, Chiba Institute of Technology, 2-17-1 Tsudanuma, Narashino, Chiba 275-0016, Japan

³High Pressure Protein Research Center, Institute of Advanced Technology, Kindai University, Kinokawa, Wakayama 649-6493, Japan

⁴Department of Mathematical and Life Sciences, School of Science, Hiroshima University, 1-3-1 Kagamiyama, Higashi-Hiroshima 739-8526, Japan

⁵Department of Physics, Nippon Medical School, 1-7-1 Kyonan-cho, Musashino, Tokyo 180-0023, Japan

⁶AMED-CREST, Japan Agency for Medical Research and Development, Japan

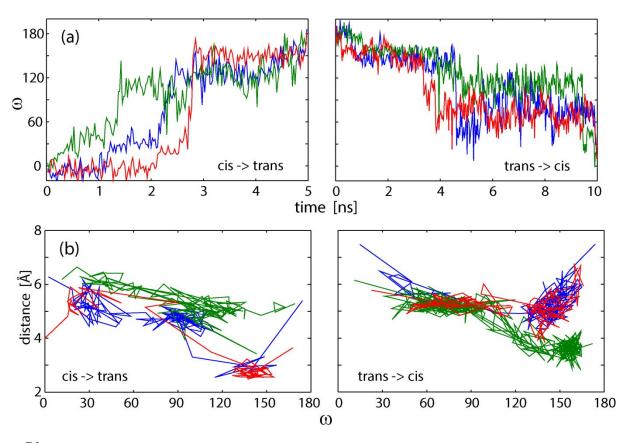


Figure S1. Three representative paths obtained from the WE simulations of both cis \rightarrow trans (left) and trans \rightarrow cis (right). (a) ω as a function of the simulation time, and (b) two-dimensional plots along ω and the atom distance between Ser154O $_{\gamma}$ and pSerO of the substrate (see Fig. 4a).

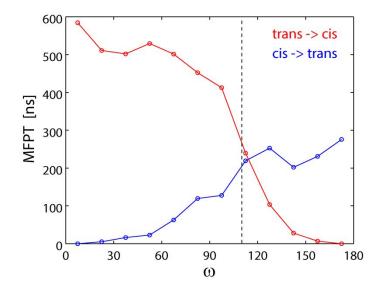


Figure S2.

Mean free passage time (MFPT) as a function of ω , calculated from the WE simulations of both cis \rightarrow trans (blue) and trans \rightarrow cis (red).