



Correlation between square of electron tunneling matrix element and donor-acceptor distance in fluctuating protein media

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Correlation between fluctuations of the square of electron tunneling matrix element TDA^2 and the donor-acceptor distance RDA in the electron transfer (ET) reaction from bacteriopheophytin anion to the primary quinone of the reaction center in the photosynthetic bacteria *Rhodospira rubra* is investigated by a combined study of molecular dynamics simulations of the protein conformation fluctuation and quantum chemical calculations. We adopted two kinds of RDA; edge-to-edge distance REE and center-to-center distance RCC. The value of TDA^2 distributed over more than 5 orders of magnitude and the fluctuation of the value of RDA distributed over more than 1.8 Å for the 106 instantaneous conformations of 1 ns simulation. We made analysis of the time-averaged correlation step by step as follows. We divide the 106 simulation data into 1000/t parts of small data set to obtain the averaged data points of t and t or t . Plotting the 1000/t sets of $\log_{10} t$ as a function of t or t , we made a principal coordinate analysis for these distributions. The slopes $\langle \beta E \rangle_t$ and $\langle \beta C \rangle_t$ of the primary axis are very large at small value of t and they are decreased considerably as t becomes large. The ellipticity for the distribution of t vs t which can be a measure for the degree of correlation became very small when t is large, while it does not hold for the distribution of t vs t . These results indicate that only the correlation between t and t for large t satisfies the well-known linear relation ("Dutton law"), although the slope is larger than the original value 1.4 \AA^{-1} . Based on the present result, we examined the analysis of the dynamic disorder by means of the single-molecule spectroscopy by Xie and co-workers with use of the "Dutton law".

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